

<210> 4093
<211> 51
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<213> Homo sapiens

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<223> 1 of 2 allelic variants (4094 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43932618

<400> 4093
aaaaacactt tggcagatgc cgtcgacaag tactgcattg gtgtgccacc c 51

<210> 4094
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4093 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43932618

<400> 4094
aaaaacactt tggcagatgc cgtcggcaag tactgcattg gtgtgccacc c 51

<210> 4095
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4096 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43932761

<400> 4095
agtttacgat cactgtcagt ttccctggag tacttaatcc gtttccttcc t 51

<210> 4096
<211> 51
<212> DNA
<213> Homo sapiens

<220>

<221> misc_feature
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<223> 2 of 2 allelic variants (4095 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43932761

<400> 4096
agtttacgat cactgtcagt ttcccaggag tacttaatcc gtttcctttc t 51

<210> 4097
<211> 51
<212> DNA
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<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4098 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43932964

<400> 4097
acttctcaca catagtaagt gggaaaagaa agtgctttga aagttcctcc c 51

<210> 4098
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4097 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43932964

<400> 4098
acttctcaca catagtaagt gggaacagaa agtgctttga aagttcctcc c 51

<210> 4099
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (4100 is other entry)

<221> misc_feature
<222> (0)...(0)

<223> Accession number cg43932964

<400> 4099

gcttcagttg gtcgaagaca gaggttcagg taaggatgac tgataggaaa t

51

<210> 4100

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 2 of 2 allelic variants (4099 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43932964

<400> 4100

gcttcagttg gtcgaagaca gaggtgcagg taaggatgac tgataggaaa t

51

<210> 4101

<211> 51

<212> DNA

<213> Homo sapiens

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<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (4102 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43933034

<400> 4101

ctgtactggtt aaaattttta cccttggtta gtctctctac tttgactaag c

51

<210> 4102

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

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<223> 2 of 2 allelic variants (4101 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43933034

<400> 4102

ctgtactggtt aaaattttta cccttcttta gtctctctac tttgactaag c

51

<210> 4103

<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4104 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43933066

<400> 4103
ataattcctc aattaatttt tattgttctt acaatatttg taagatgagt g

51

<210> 4104
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4103 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43933066

<400> 4104
ataattcctc aattaatttt tattgtcttt acaatatttg taagatgagt g

51

<210> 4105
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4106 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43933072

<400> 4105
agcgccaagt ttcttttcaa ccagtggggc ctgcagcctc gaagtctcct c

51

<210> 4106
<211> 50
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4105 is other entry)

<221> misc_feature
<222> (25)...(26)
<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43933072

<400> 4106
agcgccaagt ttcctttcaa ccagtgggcc tgcagcctcg aagtctcctc 50

<210> 4107
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4108 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43933072

<400> 4107
gccaaagtttc ctttcaacca gtggggcctg cagcctcgaa gtctcctcct c 51

<210> 4108
<211> 50
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (4107 is other entry)

<221> misc_feature
<222> (25)...(26)
<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43933072

<400> 4108
gccaaagtttc ctttcaacca gtgggcctgc agcctcgaag tctcctcctc 50

<210> 4109
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4110 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43933106

<400> 4109
caagtgtggc ggtaacgctg aggagtgagg ctgtttgtcc aggaacgctg a 51

<210> 4110
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4109 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43933106

<400> 4110
caagtgtggc ggtaacgctg aggagagagg ctgtttgtcc aggaacgctg a 51

<210> 4111
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4112 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43933106

<400> 4111
agacggtcat cagtgcagac gcagcggacg ctgctgagga tggctcagtg g 51

<210> 4112
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (4111 is other entry)

<221> misc_feature

<222> (0)...(0)
<223> Accession number cg43933106

<400> 4112
agacgggtcat cagtgcagac gcagcagacg ctgctgagga tggctcagtg g 51

<210> 4113
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4114 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43933114

<400> 4113
aaggatggct cccttccttc aacccttgat aaggggaggg aagaaaaaag a 51

<210> 4114
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4113 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43933114

<400> 4114
aaggatggct cccttccttc aaccatgat aaggggaggg aagaaaaaag a 51

<210> 4115
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4116 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43933365

<400> 4115
catctcccgc tctagaaggg ctggaagct cgcggccggg gttccacctg g 51

<210> 4116
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4115 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43933365

<400> 4116
catctccgc tctagaaggg ctggggagct cgcggccggg gttccacctg g 51

<210> 4117
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4118 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43933482

<400> 4117
ttcagctctt tttctgagat catcaccacc ccgacggaga cttgtgacga c 51

<210> 4118
<211> 51
<212> DNA
<213> Homo sapiens

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<222> (26)...(0)
<223> 2 of 2 allelic variants (4117 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43933482

<400> 4118
ttcagctctt tttctgagat catcatcacc ccgacggaga cttgtgacga c 51

<210> 4119
<211> 51
<212> DNA
<213> Homo sapiens

<220>

<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4120 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43933539

<400> 4119
cattcttgtg tagccccag tttgatcttt gtaccgaggg gccagtactt g 51

<210> 4120
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4119 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43933539

<400> 4120
cattcttgtg tagccccag tttgaccttt gtaccgaggg gccagtactt g 51

<210> 4121
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4122 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43933539

<400> 4121
ccccagtttg atctttgtac cgaggggcca gtacttgaaa cagctttggg a 51

<210> 4122
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4121 is other entry)

<221> misc_feature
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<223> Accession number cg43933539

<400> 4122

ccccagtttg atctttgtac cgaggcgcca gtacttgaaa cagctttggg a

51

<210> 4123

<211> 51

<212> DNA

<213> Homo sapiens

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<222> (26)...(0)

<223> 1 of 2 allelic variants (4124 is other entry)

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<222> (0)...(0)

<223> Accession number cg43933539

<400> 4123

tctttgtacc gaggggccag tacttgaaac agctttggga tcaaatacctt c

51

<210> 4124

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 2 of 2 allelic variants (4123 is other entry)

<221> misc_feature

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<223> Accession number cg43933539

<400> 4124

tctttgtacc gaggggccag tactttaaac agctttggga tcaaatacctt c

51

<210> 4125

<211> 51

<212> DNA

<213> Homo sapiens

<220>

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<222> (26)...(0)

<223> 1 of 2 allelic variants (4126 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43933539

<400> 4125

gtacttgaaa cagctttggg atcaaatacct tctctgtgag ccagaagtca g

51

<210> 4126

<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4125 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43933539

<400> 4126
gtacttgaaa cagctttggg atcaagtcct tctctgtgag ccagaagtca g 51

<210> 4127
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4128 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43933539

<400> 4127
acagctttgg gatcaaatcc ttctctgtga gccagaagtc agccatgcgc a 51

<210> 4128
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4127 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43933539

<400> 4128
acagctttgg gatcaaatcc ttctcagtga gccagaagtc agccatgcgc a 51

<210> 4129
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4130 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43933539

<400> 4129
tctctgtgag ccagaagtca gccatgcgca tggctcggtc gttagtgtct c 51

<210> 4130
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4129 is other entry)

<221> misc_feature
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<223> Accession number cg43933539

<400> 4130
tctctgtgag ccagaagtca gccatacgca tggctcggtc gttagtgtct c 51

<210> 4131
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4132 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43933539

<400> 4131
cagccatgcg catggctcgt tcgttagtgt ctcccagctt cccatagtcg a 51

<210> 4132
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4131 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43933539

<400> 4132
cagccatgcg catggctcgt tcgttggtgt ctcccagctt cccatagtcg a 51

<210> 4133
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4134 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43933539

<400> 4133
ctcgttcggt agtgtctccc agcttcccat agtcgatgag cttctccgcg t 51

<210> 4134
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4133 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43933539

<400> 4134
ctcgttcggt agtgtctccc agcttcccat agtcgatgag cttctccgcg t 51

<210> 4135
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4136 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43933745

<400> 4135
tcaaactcct gacctcaggt aatctgccc cctcggcctc ccaaaagtgc t 51

<210> 4136
<211> 51

<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4135 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43933745

<400> 4136
tcaaaactcct gacctcaggt aatctaccgg cctcggcctc ccaaaagtgc t 51

<210> 4137
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4138 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43933777

<400> 4137
aagattacgg ggagcagaag tctacatcca tcagcacagc aaagcgctg g 51

<210> 4138
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4137 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43933777

<400> 4138
aagattacgg ggagcagaag tctacgtcca tcagcacagc aaagcgctg g 51

<210> 4139
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)

<223> 1 of 2 allelic variants (4140 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43933777

<400> 4139

tgccattttc caagcagagc ccacggtgag gaagcacttt ctccggaaat g

51

<210> 4140

<211> 51

<212> DNA

<213> Homo sapiens

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<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (4139 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43933777

<400> 4140

tgccattttc caagcagagc ccacgatgag gaagcacttt ctccggaaat g

51

<210> 4141

<211> 51

<212> DNA

<213> Homo sapiens

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<222> (26)...(0)

<223> 1 of 2 allelic variants (4142 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43933908

<400> 4141

atttgtaata gatctgatta tatgaggtgt gaaagtcaat atgggtaatt t

51

<210> 4142

<211> 51

<212> DNA

<213> Homo sapiens

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<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (4141 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43933908

<400> 4142
atttgtaata gatctgatta tatgaagtgt gaaagtcaat atgggtaatt t 51

<210> 4143
<211> 50
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4144 is other entry)

<221> misc_feature
<222> (25)...(26)
<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43933964

<400> 4143
aaagcttttc aacctaaatg tggggaaaaa acaggtaagg cattattttt 50

<210> 4144
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4143 is other entry)

<221> misc_feature
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<223> Accession number cg43933964

<400> 4144
aaagcttttc aacctaaatg tggggaaaaa aacaggtaag gcattatttt t 51

<210> 4145
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4146 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43934149

<400> 4145
tatgttatct attcagtttt gaaaacattc attaagattt taaatgcaaa t 51

<210> 4146
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4145 is other entry)

<221> misc_feature
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<223> Accession number cg43934149

<400> 4146
tatgttatct attcagtttt gaaaaaatc attaagattt taaatgcaa t 51

<210> 4147
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4148 is other entry)

<221> misc_feature
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<223> Accession number cg43934157

<400> 4147
ttaaccatta ttctataaga cataaggga ggtaaataat ggcccacaaa a 51

<210> 4148
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4147 is other entry)

<221> misc_feature
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<223> Accession number cg43934157

<400> 4148
ttaaccatta ttctataaga cataaggga ggtaaataat ggcccacaaa a 51

<210> 4149
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4150 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43934300

<400> 4149
ccaggatagt taacttgaat ttcacagtg tccacagtca gaggattaaa g 51

<210> 4150
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4149 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43934300

<400> 4150
ccaggatagt taacttgaat ttcataagtg tccacagtca gaggattaaa g 51

<210> 4151
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4152 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43934316

<400> 4151
cattaacaaa acaattctgg tactacagac cagtggtgac agaattaggct t 51

<210> 4152
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 2 of 2 allelic variants (4151 is other entry)

<221> misc_feature

<222> (0)...(0)
<223> Accession number cg43934316

<400> 4152
cattaacaaa acaattctgg tactagagac cagtgggtgc agaataggct t 51

<210> 4153
<211> 51
<212> DNA
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<223> 1 of 2 allelic variants (4154 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43934316

<400> 4153
aaacaattct ggtactacag accagtgggtg tcagaatagg cttagtgcct c 51

<210> 4154
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (4153 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43934316

<400> 4154
aaacaattct ggtactacag accagcgggtg tcagaatagg cttagtgcct c 51

<210> 4155
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4156 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43934316

<400> 4155
gtactacaga ccagtgggtg cagaataggc ttagtgcttc cttgtttgtg t 51

<210> 4156
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4155 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43934316

<400> 4156
gtactacaga ccagtgggtg cagaacaggc ttagtgccctc cttgtttggtg t 51

<210> 4157
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4158 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43934316

<400> 4157
tcagagttca ttttatgcag ggccattctc agtcctcaat gtactccac a 51

<210> 4158
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4157 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43934316

<400> 4158
tcagagttca ttttatgcag ggccagtctc agtcctcaat gtactccac a 51

<210> 4159
<211> 51
<212> DNA
<213> Homo sapiens

<220>

<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4160 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43934316

<400> 4159
cagagttcat tttatgcagg gccattctca gtctcaatg tactcccaca g 51

<210> 4160
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (4159 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43934316

<400> 4160
cagagttcat tttatgcagg gccatgctca gtctcaatg tactcccaca g 51

<210> 4161
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4162 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43934316

<400> 4161
actgcaatac cctcaggcag tatgccaaca ttgaaataga aagcatctct a 51

<210> 4162
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4161 is other entry)

<221> misc_feature
<222> (0)...(0)

<223> Accession number cg43934316

<400> 4162
actgcaatac cctcaggcag tatgctaaca ttgaaataga aagcatctct a 51

<210> 4163
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4164 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43934439

<400> 4163
ggatgatgca ggtatggagt tgcagcccc acagcagaca ttgctgctgc t 51

<210> 4164
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4163 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43934439

<400> 4164
ggatgatgca ggtatggagt tgcagtcccc acagcagaca ttgctgctgc t 51

<210> 4165
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4166 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43934462

<400> 4165
ggaatctgtg acccgcttg tgactggtgg gcagattcaa gtctctaccg c 51

<210> 4166

<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4165 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43934462

<400> 4166
ggaatctgtg acccgcttg tgactagtgg gcagattcaa gtctctaccg c 51

<210> 4167
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4168 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43934499

<400> 4167
tgtgtcttcc tgtgttctca agattgagaa tcccagatga tcacgcgatt t 51

<210> 4168
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4167 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43934499

<400> 4168
tgtgtcttcc tgtgttctca agatttagaa tcccagatga tcacgcgatt t 51

<210> 4169
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature

<222> (26)...(0)
<223> 1 of 2 allelic variants (4170 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43934499

<400> 4169
tctcaagatt gagaatccca gatgatcacg cgatttcaga cctatccatg t 51

<210> 4170
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4169 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43934499

<400> 4170
tctcaagatt gagaatccca gatgaccacg cgatttcaga cctatccatg t 51

<210> 4171
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4172 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43934499

<400> 4171
atgatcacgc gatttcagac ctatccatgt acaattacat agagatgcga g 51

<210> 4172
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4171 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43934499

<400> 4172
atgatcacgc gatttcagac ctatctatgt acaattacat agagatgcga g 51

<210> 4173
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4174 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43934499

<400> 4173
tgtacaatta catagagatg cgagcacatg tcaactcaag ctgggttcatt t 51

<210> 4174
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4173 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43934499

<400> 4174
tgtacaatta catagagatg cgagcgcgatg tcaactcaag ctgggttcatt t 51

<210> 4175
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4176 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43934499

<400> 4175
tacatagaga tgcgagcaca tgtcaactca agctgggttca tttttcagaa g 51

<210> 4176
<211> 51

<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4175 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43934499

<400> 4176
tacatagaga tgcgagcaca tgtcatctca agctgggttca tttttcagaa g 51

<210> 4177
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (4178 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43934499

<400> 4177
catcgacctt tatccctctc tatacaatgg tcactttttc cagaataaga t 51

<210> 4178
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4177 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43934499

<400> 4178
catcgacctt tatccctctc tataccatgg tcactttttc cagaataaga t 51

<210> 4179
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (4180 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43934665

<400> 4179

cccgggttcg agcgattttc ctgcctcagc ctcccaagta gggggactac a

51

<210> 4180

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (4179 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43934665

<400> 4180

cccgggttcg agcgattttc ctgccccagc ctcccaagta gggggactac a

51

<210> 4181

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (4182 is other entry)

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<222> (0)...(0)

<223> Accession number cg43934665

<400> 4181

gcctcagcct cccaagtagg gggactacag gcaccaccca ccacgccgg c

51

<210> 4182

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 2 of 2 allelic variants (4181 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43934665

<400> 4182
gcctcagcct cccaagtagg gggaccacag gcacccacca ccacgcccg c 51

<210> 4183
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4184 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43935007

<400> 4183
gtggagattg tagaataact atcattagca aaggcagaaa gtattcattt c 51

<210> 4184
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4183 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43935007

<400> 4184
gtggagattg tagaataact atcatgagca aaggcagaaa gtattcattt c 51

<210> 4185
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4186 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43935063

<400> 4185
ggcaccagga agtccacacc atatgcagta ttggggctgt aggtctccga g 51

<210> 4186
<211> 51
<212> DNA

<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4185 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43935063

<400> 4186
ggcaccagga agtccacacc atatggagta ttggggctgt aggtctccga g 51

<210> 4187
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4188 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43935145

<400> 4187
cctccatcac cacgaacatg tggtaggccg gcgcgggccc ccgctccaag a 51

<210> 4188
<211> 50
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4187 is other entry)

<221> misc_feature
<222> (25)...(26)
<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43935145

<400> 4188
cctccatcac cacgaacatg tggtagccgc gccgggcccc cgctccaaga 50

<210> 4189
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4190 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43935384

<400> 4189
ttgcagattt ttcttctaaa aaaaaactat aattctctca cagatcacat a 51

<210> 4190
<211> 50
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4189 is other entry)

<221> misc_feature
<222> (25)...(26)
<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43935384

<400> 4190
ttgcagattt ttcttctaaa aaaaactata attctctcac agatcacata 50

<210> 4191
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4192 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43935526

<400> 4191
tttttttcca aaatcactgt tgggggtgggg gatccagtc tcgggactgt g 51

<210> 4192
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4191 is other entry)

<221> misc_feature
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<223> Accession number cg43935526

<400> 4192
tttttttcca aaatcactgt tggggggggg gatcccagtc tcgggactgt g 51

<210> 4193
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4194 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43935836

<400> 4193
tggccttggc cagacacaaa ccaagagact gccatgacag acagagcaga a 51

<210> 4194
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4193 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43935836

<400> 4194
tggccttggc cagacacaaa ccaagggact gccatgacag acagagcaga a 51

<210> 4195
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4196 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43935861

<400> 4195
tggcagagaa aaagggccacc gatgctgaag cgcacgtagc ttctctgaac a 51

<210> 4196
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4195 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43935861

<400> 4196
tggcagagaa aaagggccacc gatgcggaag cgcacgtagc ttctctgaac a 51

<210> 4197
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4198 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43935933

<400> 4197
atgatgacac tgtgcctgcc aagggcagtt tgcttttctt caactctagg c 51

<210> 4198
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4197 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43935933

<400> 4198
atgatgacac tgtgcctgcc aaggctagtt tgcttttctt caactctagg c 51

<210> 4199
<211> 51

<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4200 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43936041

<400> 4199
gtggaaaagc agatctgtgg atgtcaagcc gaagatcact ccgtttatgg a 51

<210> 4200
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4199 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43936041

<400> 4200
gtggaaaagc agatctgtgg atgtcgagcc gaagatcact ccgtttatgg a 51

<210> 4201
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4202 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43936051

<400> 4201
tgttaccagc ttacatact gttctgccat ttgtgagggg tgcaaccaca a 51

<210> 4202
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)

<223> 2 of 2 allelic variants (4201 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43936051

<400> 4202

tgttaccagc ttacatact gttctacat ttgtgagggg tgcaaccaca a

51

<210> 4203

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (4204 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43936117

<400> 4203

ggagggcgag cagaaaaacg gaaaacacgg aacgccacag aagtatgatc c

51

<210> 4204

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (4203 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43936117

<400> 4204

ggagggcgag cagaaaaacg gaaaatacgg aacgccacag aagtatgatc c

51

<210> 4205

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (4206 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43936154

<400> 4205
ggatgccgtg aggggatgtg atgtcctcag tgcctctgat gacacagttg c 51

<210> 4206
<211> 50
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4205 is other entry)

<221> misc_feature
<222> (25)...(26)
<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43936154

<400> 4206
ggatgccgtg aggggatgtg atgtctcagt gcctctgatg acacagttgc 50

<210> 4207
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4208 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43936249

<400> 4207
ccacgatecc ttctactcat tggtagcac accagattag gtacaagaat c 51

<210> 4208
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 2 of 2 allelic variants (4207 is other entry)

<221> misc_feature
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<223> Accession number cg43936249

<400> 4208
ccacgatecc ttctactcat tggtaggcac accagattag gtacaagaat c 51

<210> 4209
<211> 51
<212> DNA
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<223> 1 of 2 allelic variants (4210 is other entry)

<221> misc_feature
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<223> Accession number cg43936432

<400> 4209
tatgctgcct gaaatggcct atgcctccta aatttccttt cactttgtca c 51

<210> 4210
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 2 of 2 allelic variants (4209 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43936432

<400> 4210
tatgctgcct gaaatggcct atgcccccta aatttccttt cactttgtca c 51

<210> 4211
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (4212 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43936887

<400> 4211
tctgttgctc cttcagtcag gtcactgttc aaatagctct ctagacaggc t 51

<210> 4212
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 2 of 2 allelic variants (4211 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43936887

<400> 4212
tctgttggtcc cttcagtcag gtcaccgttc aaatagctct ctagacaggc t 51

<210> 4213
<211> 51
<212> DNA
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<223> 1 of 2 allelic variants (4214 is other entry)

<221> misc_feature
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<223> Accession number cg43936887

<400> 4213
cttccttatc attctactta aaatagcccc caatcactct gtgtcccttt a 51

<210> 4214
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 2 of 2 allelic variants (4213 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43936887

<400> 4214
cttccttatc attctactta aaataacccc caatcactct gtgtcccttt a 51

<210> 4215
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (4216 is other entry)

<221> misc_feature

<222> (0)...(0)
<223> Accession number cg43936887

<400> 4215
tcttctgct atttcatact acctgaaaaa aatacttgaa cttcctagaa c 51

<210> 4216
<211> 50
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4215 is other entry)

<221> misc_feature
<222> (25)...(26)
<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43936887

<400> 4216
tcttctgct atttcatact acctgaaaaa atacttgaac ttcctagaac 50

<210> 4217
<211> 51
<212> DNA
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<220>
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<223> 1 of 2 allelic variants (4218 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43936887

<400> 4217
tgctatttca tactacctga aaaaaaatact tgaacttcct agaacataag c 51

<210> 4218
<211> 50
<212> DNA
<213> Homo sapiens

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<223> 2 of 2 allelic variants (4217 is other entry)

<221> misc_feature
<222> (25)...(26)
<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43936887

<400> 4218
tgctattttca tactacctga aaaaatactt gaacttccta gaacataagc 50

<210> 4219
<211> 51
<212> DNA
<213> Homo sapiens

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<222> (26)...(0)
<223> 1 of 2 allelic variants (4220 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43936905

<400> 4219
agtttttctg ccgctggcag ggctgcgggg accgccagct gctgcagtgc g 51

<210> 4220
<211> 50
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (4219 is other entry)

<221> misc_feature
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<221> misc_feature
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<223> Accession number cg43936905

<400> 4220
agtttttctg ccgctggcag ggctggggga ccgccagctg ctgcagtgcg 50

<210> 4221
<211> 51
<212> DNA
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4222 is other entry)

<221> misc_feature

<222> (0)...(0)
<223> Accession number cg43937728

<400> 4221
ggctttgtca cggtgagcta cttctcacag gagtcgggat ccacatctgc a 51

<210> 4222
<211> 51
<212> DNA
<213> Homo sapiens

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<222> (26)...(0)
<223> 2 of 2 allelic variants (4221 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43937728

<400> 4222
ggctttgtca cggtgagcta cttcttacag gagtcgggat ccacatctgc a 51

<210> 4223
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 1 of 2 allelic variants (4224 is other entry)

<221> misc_feature
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<223> Accession number cg43937732

<400> 4223
gttgccccac aggggaaggg gcgcccgggc gcggccgccg gaggcatttg g 51

<210> 4224
<211> 51
<212> DNA
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<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4223 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43937732

<400> 4224
gttgccccac aggggaaggg gcgcctgggc gcggccgccg gaggcatttg g 51

<210> 4225

<211> 50

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (4226 is other entry)

<221> misc_feature

<222> (25)...(26)

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<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43937732

<400> 4225

tatttttgga ttaccaatt tttttctac tattctcaga tatctatcaa

50

<210> 4226

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (4225 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43937732

<400> 4226

tatttttgga ttaccaatt ttttttccta ctattctcag atatctatca a

51

<210> 4227

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (4228 is other entry)

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<222> (0)...(0)

<223> Accession number cg43937732

<400> 4227

cttaaggcac agatcactca tgctattggt tgtggtttag gaatgccttt a

51

<210> 4228

<211> 51

<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4227 is other entry)

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<223> Accession number cg43937732

<400> 4228
cttaaggcac agatcactca tgctactggt tgggttttag gaatgccttt a 51

<210> 4229
<211> 51
<212> DNA
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<223> 1 of 2 allelic variants (4230 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43937732

<400> 4229
ctgggtgggc caggtgttcc ttgcctcat tccggtaaac ccaaacctt c 51

<210> 4230
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4229 is other entry)

<221> misc_feature
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<223> Accession number cg43937732

<400> 4230
ctgggtgggc caggtgttcc ttgcctcat tccggtaaac ccaaacctt c 51

<210> 4231
<211> 51
<212> DNA
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<223> 1 of 2 allelic variants (4232 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43937732

<400> 4231

gggccttctc ccaatatgtc ccccttcttt gatttgcaaa tcgataaaag c

51

<210> 4232

<211> 51

<212> DNA

<213> Homo sapiens

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<222> (26)...(0)

<223> 2 of 2 allelic variants (4231 is other entry)

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<222> (0)...(0)

<223> Accession number cg43937732

<400> 4232

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<210> 4233

<211> 51

<212> DNA

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<223> 1 of 2 allelic variants (4234 is other entry)

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<222> (0)...(0)

<223> Accession number cg43938515

<400> 4233

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51

<210> 4234

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 2 of 2 allelic variants (4233 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43938515

<400> 4234
agatgggggg tgatttcagc atcacacacc ctccatatatg gccaggcctc c 51

<210> 4235
<211> 51
<212> DNA
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<220>
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<223> 1 of 2 allelic variants (4236 is other entry)

<221> misc_feature
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<223> Accession number cg43938812

<400> 4235
cactgacagt gcccccggtg cgtgcatgta ttctgcgcac tttcctgtgc t 51

<210> 4236
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (4235 is other entry)

<221> misc_feature
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<223> Accession number cg43938812

<400> 4236
cactgacagt gcccccggtg cgtgcgtgta ttctgcgcac tttcctgtgc t 51

<210> 4237
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4238 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43938812

<400> 4237
ctgcctctgt gcctgcctgt actgccgatg ctccagtggg taactcagca t 51

<210> 4238
<211> 51
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<213> Homo sapiens

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<222> (26)...(0)

<223> 2 of 2 allelic variants (4237 is other entry)

<221> misc_feature

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<223> Accession number cg43938812

<400> 4238

ctgcctctgt gctgcctgt actgctgatg ctccagtga taactcagca t

51

<210> 4239

<211> 51

<212> DNA

<213> Homo sapiens

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<222> (26)...(0)

<223> 1 of 2 allelic variants (4240 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43939553

<400> 4239

aagaacccac aagtgtccag agggatttct agtgatctc tctcttaacc c

51

<210> 4240

<211> 50

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (4239 is other entry)

<221> misc_feature

<222> (25)...(26)

<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43939553

<400> 4240

aagaacccac aagtgtccag agggatttcta ggtgatctct ctcttaaccc

50

<210> 4241

<211> 51

<212> DNA

<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4242 is other entry)

<221> misc_feature
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<223> Accession number cg43939553

<400> 4241
gaaccacaa gtgtccagag ggatttctag gtgatctctc tcttaacccc t 51

<210> 4242
<211> 50
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4241 is other entry)

<221> misc_feature
<222> (25)...(26)
<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43939553

<400> 4242
gaaccacaa gtgtccagag ggattctagg tgatctctct cttaacccct 50

<210> 4243
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4244 is other entry)

<221> misc_feature
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<223> Accession number cg43939553

<400> 4243
aaaagaagaa gaaactcaaa attcctatct gcgtgctaatttgaaaagaa c 51

<210> 4244
<211> 51
<212> DNA
<213> Homo sapiens

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<222> (26)...(0)
<223> 2 of 2 allelic variants (4243 is other entry)

<221> misc_feature
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<400> 4244
aaaagaagaa gaaactcaaa attcccatct gcgtgctaatttgaaaagaa c 51

<210> 4245
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4246 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43939553

<400> 4245
atcaagagca aagggaacag caggcctaac agcagggttg ggaaggcaaa a 51

<210> 4246
<211> 50
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4245 is other entry)

<221> misc_feature
<222> (25)...(26)
<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43939553

<400> 4246
atcaagagca aagggaacag caggctaaca gcagggttgg gaaggcaaaa 50

<210> 4247
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4248 is other entry)

<221> misc_feature
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<223> Accession number cg43939553

<400> 4247
tagcagggcat ttataagtgc ccaccctcac caatgcatcg ggggtgggtcc c 51

<210> 4248
<211> 50
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4247 is other entry)

<221> misc_feature
<222> (25)...(26)
<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43939553

<400> 4248
tagcagggcat ttataagtgc ccacctcacc aatgcatcgg ggggtgggtccc 50

<210> 4249
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4250 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43940188

<400> 4249
gctgctggcc aaggcgggagc gcgtgagctc gcacgccaac gccgccaag a 51

<210> 4250
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
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<223> 2 of 2 allelic variants (4249 is other entry)

<221> misc_feature

<222> (0)...(0)
<223> Accession number cg43940188

<400> 4250
gctgctggcc aaggcggagc gcgtgggctc gcacgccaac gccgcccaag a 51

<210> 4251
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4252 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43940467

<400> 4251
agtacataaa taaatactaa aaaaaattaa aatccttggt cttattttgt a 51

<210> 4252
<211> 50
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4251 is other entry)

<221> misc_feature
<222> (25)...(26)
<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43940467

<400> 4252
agtacataaa taaatactaa aaaaattaaa atccttggtc ttattttgta 50

<210> 4253
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4254 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43941368

<400> 4253
ggcagaatcc tgctattccc aagaaccctc gtaatggcaa aactccccaa a 51

<210> 4254
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4253 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43941368

<400> 4254
ggcagaatcc tgctattccc aagaagcctc gtaatggcaa aactccccaa a 51

<210> 4255
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4256 is other entry)

<221> misc_feature
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<223> Accession number cg43941368

<400> 4255
atcctgctat tccaagaac cctcgtaatg gcaaaactcc ccaaagaca c 51

<210> 4256
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4255 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43941368

<400> 4256
atcctgctat tccaagaac cctcgcaatg gcaaaactcc ccaaagaca c 51

<210> 4257
<211> 51

<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4258 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43941368

<400> 4257
cccaagaacc ctcgtaatgg caaaactccc caaatgacac ccaggaccac a 51

<210> 4258
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4257 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43941368

<400> 4258
cccaagaacc ctcgtaatgg caaaattccc caaatgacac ccaggaccac a 51

<210> 4259
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4260 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43941368

<400> 4259
cccaaagac acccaggacc acagcaatga tctgtcggaa ccagtagatc a 51

<210> 4260
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)

<223> 2 of 2 allelic variants (4259 is other entry)

<221> misc_feature
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<223> Accession number cg43941368

<400> 4260
cccaaagac acccaggacc acagcgatga tctgtcggaa ccagtagatc a 51

<210> 4261
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4262 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43941470

<400> 4261
agagctgctg tagtctctgc ttggcctctt tgctcagctt caccatggcg a 51

<210> 4262
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4261 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43941470

<400> 4262
agagctgctg tagtctctgc ttggcctctt tgctcagctt caccatggcg a 51

<210> 4263
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4264 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43941536

<400> 4263
ttctctgtag ctctgaaaaa caggaacagg cattgaacag ggccttgacc a 51

<210> 4264
<211> 50
<212> DNA
<213> Homo sapiens

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<222> (26)...(0)
<223> 2 of 2 allelic variants (4263 is other entry)

<221> misc_feature
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<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43941536

<400> 4264
ttctctgtag ctctgaaaaa caggacaggc attgaacagg gccttgacca 50

<210> 4265
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4266 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43941567

<400> 4265
cctccgcct cggcctccca aagtgtggg attacaggcg tgagccaccg c 51

<210> 4266
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (4265 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43941567

<400> 4266
cctccgcct cggcctccca aagtgtggg attacaggcg tgagccaccg c 51

<210> 4267
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4268 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43942501

<400> 4267
cctttctcta ggcggctctc tggctctgac ttcacagcc agtggctgct t 51

<210> 4268
<211> 51
<212> DNA
<213> Homo sapiens

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<222> (26)...(0)
<223> 2 of 2 allelic variants (4267 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43942501

<400> 4268
cctttctcta ggcggctctc tggctttgac ttcacagcc agtggctgct t 51

<210> 4269
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4270 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43942588

<400> 4269
atgaaatcat ggagcagaag tccaggagag ttcaatacga tttctaaatc c 51

<210> 4270
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4269 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43942588

<400> 4270
atgaaatcat ggagcagaag tccagtagag ttcaatacga tttctaaatc c 51

<210> 4271
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4272 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43942890

<400> 4271
ttgctcactc cgaggtggaa taaggagac gttagtagtc acatatgtac t 51

<210> 4272
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (4271 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43942890

<400> 4272
ttgctcactc cgaggtggaa taaggaagac gttagtagtc acatatgtac t 51

<210> 4273
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4274 is other entry)

<221> misc_feature

<222> (0)...(0)
<223> Accession number cg43943086

<400> 4273
tctagctatt ttcaagacag acttaatcaa tactgtgttt gctttccaat t 51

<210> 4274
<211> 50
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (4273 is other entry)

<221> misc_feature
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<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43943086

<400> 4274
tctagctatt ttcaagacag acttatcaat actgtgtttg ctttccaatt 50

<210> 4275
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4276 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43943163

<400> 4275
tggttctcgg cctcctgttt ctgcacatgg cgccgcaggg tgtcccgttg c 51

<210> 4276
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4275 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43943163

<400> 4276
tggttctcgg cctcctgttt ctgcatatgg cgccgcaggg tgccccgttg c 51

<210> 4277
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4278 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43943237

<400> 4277
catcttgaga agaaactaac ttctgccttt aatttgcata taagtatcat a 51

<210> 4278
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4277 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43943237

<400> 4278
catcttgaga agaaactaac ttctgtcttt aatttgcata taagtatcat a 51

<210> 4279
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4280 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43943237

<400> 4279
cttaactgtt accactatag tcaagcccag tacctcagtg acttcacaga t 51

<210> 4280
<211> 51

<212> DNA
<213> Homo sapiens

<220>
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<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43943237

<400> 4280
cttaactggt accactatag tcaagtccag tacctcagtg acttcacaga t 51

<210> 4281
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4282 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43943237

<400> 4281
aactttaagt tcctatgaca tagtacggtg ttagtatggt gtatagctgt a 51

<210> 4282
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4281 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43943237

<400> 4282
aactttaagt tcctatgaca tagtatggtg ttagtatggt gtatagctgt a 51

<210> 4283
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)

<223> 1 of 2 allelic variants (4284 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43943237

<400> 4283

ggagttttgc tactaggttt tttttgttt ttgtttttt ttacaaatc a

51

<210> 4284

<211> 50

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (4283 is other entry)

<221> misc_feature

<222> (25)...(26)

<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43943237

<400> 4284

ggagttttgc tactaggttt tttttgttt ttgtttttt ttacaaatca

50

<210> 4285

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (4286 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43943787

<400> 4285

aaagtccgaa atcactgac ttggcgtaat gttgggtaac tagcaacaca t

51

<210> 4286

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (4285 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43943787 ,

<400> 4286
aaagtccgaa atcactgatc ttggcataat gttgggtaac tagcaacaca t 51

<210> 4287
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4288 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43944032

<400> 4287
tgacagaagc tcattaaaac caagtgcgcc aaacctcctg aaacatcggt a 51

<210> 4288
<211> 51
<212> DNA
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<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4287 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43944032

<400> 4288
tgacagaagc tcattaaaac caagtgcgcc aaacctcctg aaacatcggt a 51

<210> 4289
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4290 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43944291

<400> 4289
ccacttctga acggccgaag gtgccccatt ccagacctgc ccatttgatg g 51

<210> 4290
<211> 51
<212> DNA
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<223> 2 of 2 allelic variants (4289 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43944291

<400> 4290
ccacttctga acggccgaag gtgcctcatt ccagacctgc ccatttgatg g 51

<210> 4291
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4292 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43944291

<400> 4291
atatttcttc acaggctgtg gaattctcta gctaaacatt ctagtttctc c 51

<210> 4292
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4291 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43944291

<400> 4292
atatttcttc acaggctgtg gaatttctcta gctaaacatt ctagtttctc c 51

<210> 4293
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4294 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43944291

<400> 4293
taaaaaatac attcatacag aaatataaca atcttgcaaa aaacaatttc a 51

<210> 4294
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4293 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43944291

<400> 4294
taaaaaatac attcatacag aaataaaaaca atcttgcaaa aaacaatttc a 51

<210> 4295
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4296 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43944291

<400> 4295
agggtgcttc caaaaaaaaaa aaaaaagaaa tttcactaat agaaattttt t 51

<210> 4296
<211> 50
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4295 is other entry)

<221> misc_feature

<222> (25)...(26)
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<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43944291

<400> 4296
agggtgcttc caaaaaaaaa aaaaagaaat ttcactaata gaaatttttt 50

<210> 4297
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4298 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43944310

<400> 4297
ggtatttaac tggaataag ggagacacaa accatttctc tcccgcactc c 51

<210> 4298
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4297 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43944310

<400> 4298
ggtatttaac tggaataag ggagaaacaa accatttctc tcccgcactc c 51

<210> 4299
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4300 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43944408

<400> 4299
ttgttgtag tcattgtaga tggatcgtct ggaattccta gaggaagagg a 51

<210> 4300
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4299 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43944408

<400> 4300
ttgttgtag tcattgtaga tggatcgtct ggaattccta gaggaagagg a 51

<210> 4301
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4302 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43944408

<400> 4301
agccggcatc tcttaacttt ttttttcccc ccagtaaatt ggtatgcaat a 51

<210> 4302
<211> 50
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4301 is other entry)

<221> misc_feature
<222> (25)...(26)
<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43944408

<400> 4302

agccggcatc tcttaacttt tttttccccc cagtaaattg gtatgcaata

50

<210> 4303

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (4304 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43944446

<400> 4303

actgggacca atccagtggg cagcgccctc ccagggtac attccagaac a

51

<210> 4304

<211> 50

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (4303 is other entry)

<221> misc_feature

<222> (25)...(26)

<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43944446

<400> 4304

actgggacca atccagtggg cagcgccctc ccagggtaca ttccagaaca

50

<210> 4305

<211> 50

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (4306 is other entry)

<221> misc_feature

<222> (25)...(26)

<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43944446

<400> 4305
atcgggttatt atccccctcat ttttttgtagg aaataagttt gcttgtttct 50

<210> 4306
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4305 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43944446

<400> 4306
atcgggttatt atccccctcat ttttttgtag gaaataagtt tgcttgtttc t 51

<210> 4307
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4308 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43945296

<400> 4307
ggggcagtg gaaaccgccac cggggccgct gtagcgggcc ttaaaggatg g 51

<210> 4308
<211> 50
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4307 is other entry)

<221> misc_feature
<222> (25)...(26)
<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43945296

<400> 4308

ggggcagtgg gaaccgccac cggggcgctg tagcgggcct taaaggatgg

50

<210> 4309

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (4310 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43945296

<400> 4309

gggcagtggg aaccgccacc ggggcccgtg tagcgggcct taaaggatgg g

51

<210> 4310

<211> 50

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

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<223> 2 of 2 allelic variants (4309 is other entry)

<221> misc_feature

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<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43945296

<400> 4310

gggcagtggg aaccgccacc ggggcccgtgt agcgggcctt aaaggatggg

50

<210> 4311

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

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<223> 1 of 2 allelic variants (4312 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43945296

<400> 4311

gcgggcctta aaggatggga aaccttgatc acagatgcc cccgccggcc t

51

<210> 4312
<211> 50
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4311 is other entry)

<221> misc_feature
<222> (25)...(26)
<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43945296

<400> 4312
gcgggcctta aaggatggga aacctgatca cagatccccc ccgccggcct

50

<210> 4313
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4314 is other entry)

<221> misc_feature
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<223> Accession number cg43945296

<400> 4313
ctccccgagc gcaggccctt ctctctttgc cttattttatt tttggaacat a

51

<210> 4314
<211> 51
<212> DNA
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<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4313 is other entry)

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<223> Accession number cg43945296

<400> 4314
ctccccgagc gcaggccctt ctctccttgc cttattttatt tttggaacat a

51

<210> 4315
<211> 51

<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4316 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43946435

<400> 4315
gggctgctgg tcccaaccag ctggtgctg tggtggatg gttcagtgt g 51

<210> 4316
<211> 50
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4315 is other entry)

<221> misc_feature
<222> (25)...(26)
<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43946435

<400> 4316
gggctgctgg tcccaaccag ctggtcctgt ggctggatgt gttcagtgtg 50

<210> 4317
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4318 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43946684

<400> 4317
aaaacaacaa taaaattctc tttgaaggga accaaagaca atgatgtgtt c 51

<210> 4318
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4317 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43946684

<400> 4318
aaaacaacaa taaaattctc ttgacggga accaaagaca atgatgtgtt c 51

<210> 4319
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4320 is other entry)

<221> misc_feature
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<223> Accession number cg43946814

<400> 4319
cccaggaagg tggcgccccg ctccccagcc tgctacaggg aaccgggac t 51

<210> 4320
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (4319 is other entry)

<221> misc_feature
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<223> Accession number cg43946814

<400> 4320
cccaggaagg tggcgccccg ctccaagcc tgctacaggg aaccgggac t 51

<210> 4321
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (4322 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43947005

<400> 4321
aactgtacaa gcaattaaaa catgatatgt agcaagtgtt atcaggagtt t 51

<210> 4322
<211> 51
<212> DNA
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<220>
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<223> 2 of 2 allelic variants (4321 is other entry)

<221> misc_feature
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<223> Accession number cg43947005

<400> 4322
aactgtacaa gcaattaaaa catgaaatgt agcaagtgtt atcaggagtt t 51

<210> 4323
<211> 51
<212> DNA
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<220>
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<223> 1 of 2 allelic variants (4324 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43947375

<400> 4323
ggcgatctcc acatcctgcc ggtggcattt aggggtgact ccttcacaca t 51

<210> 4324
<211> 50
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (4323 is other entry)

<221> misc_feature
<222> (25)...(26)
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<221> misc_feature
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<223> Accession number cg43947375

<400> 4324

ggcgatctcc acatcctgcc ggtggattta ggggtgactc cttcacacat

50

<210> 4325

<211> 51

<212> DNA

<213> Homo sapiens

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<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (4326 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43947477

<400> 4325

acaaatttgg cttaataaa aaaaaaacac gttcaaaagg acaataacac g

51

<210> 4326

<211> 50

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (4325 is other entry)

<221> misc_feature

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<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43947477

<400> 4326

acaaatttgg cttaataaa aaaaaaacag ttcaaaagga caataacacg

50

<210> 4327

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (4328 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43947477

<400> 4327
caaatttggc ttttaataaaa aaaaaaacag ttcaaaagga caataacacg g 51

<210> 4328
<211> 50
<212> DNA
<213> Homo sapiens

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<223> 2 of 2 allelic variants (4327 is other entry)

<221> misc_feature
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<221> misc_feature
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<223> Accession number cg43947477

<400> 4328
caaatttggc ttttaataaaa aaaaaacagt tcaaaaggac aataacacgg 50

<210> 4329
<211> 51
<212> DNA
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<220>
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<223> 1 of 2 allelic variants (4330 is other entry)

<221> misc_feature
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<223> Accession number cg43947477

<400> 4329
aaatttggct ttaataaaaa aaaaaacagt tcaaaaggac aataacacgg g 51

<210> 4330
<211> 50
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (4329 is other entry)

<221> misc_feature
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<221> misc_feature
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<223> Accession number cg43947477

<400> 4330

aaatttggt ttaataaaaa aaaaacagtt caaaaggaca ataacacggg

50

<210> 4331

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

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<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43947477

<400> 4331

tgtgcagttt cctgatcata atcacatgtc tectgccttt tacagggaat g

51

<210> 4332

<211> 50

<212> DNA

<213> Homo sapiens

<220>

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<223> 2 of 2 allelic variants (4331 is other entry)

<221> misc_feature

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<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43947477

<400> 4332

tgtgcagttt cctgatcata atcactgtct cctgcctttt acagggaatg

50

<210> 4333

<211> 51

<212> DNA

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<223> 1 of 2 allelic variants (4334 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43947477

<400> 4333
cccattctcc tatggaatgg ttttcagggg tccagcaatg tcaactgcagg t 51

<210> 4334
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (4333 is other entry)

<221> misc_feature
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<223> Accession number cg43947477

<400> 4334
cccattctcc tatggaatgg ttttcagggg tccagcaatg tcaactgcagg t 51

<210> 4335
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4336 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43947477

<400> 4335
gggagactct gttgtaggca gctgtgggag aaggctaggg ttggaggtac a 51

<210> 4336
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (4335 is other entry)

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<223> Accession number cg43947477

<400> 4336
gggagactct gttgtaggca gctgtcggag aaggctaggg ttggaggtac a 51

<210> 4337
<211> 51
<212> DNA

<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4338 is other entry)

<221> misc_feature
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<223> Accession number cg43947638

<400> 4337
agatttgagt gtgcgaggaa aaaataaaaa agagaaactt gaagacttct t 51

<210> 4338
<211> 50
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4337 is other entry)

<221> misc_feature
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<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43947638

<400> 4338
agatttgagt gtgcgaggaa aaaataaaaa gagaaacttg aagacttctt 50

<210> 4339
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4340 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43947646

<400> 4339
ctcttcacat cctctatatt ctgtgcgact tgaaagctgt ttgagaactt c 51

<210> 4340
<211> 51
<212> DNA
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<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4339 is other entry)

<221> misc_feature
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<223> Accession number cg43947646

<400> 4340
ctcttcacat cctctatatt ctgtgtgact tgaaagctgt ttgagaactt c 51

<210> 4341
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4342 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43947743

<400> 4341
acacttttgg gaagcctggg accatggctc tgccaggaat ctgtgacatc t 51

<210> 4342
<211> 50
<212> DNA
<213> Homo sapiens

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<222> (26)...(0)
<223> 2 of 2 allelic variants (4341 is other entry)

<221> misc_feature
<222> (25)...(26)
<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43947743

<400> 4342
acacttttgg gaagcctggg accatgctct gccaggaatc tgtgacatct 50

<210> 4343
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4344 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43947977

<400> 4343
gagcaggacc tccattagaa atattgaatt tgattccaaa atctccattg g 51

<210> 4344
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 2 of 2 allelic variants (4343 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43947977

<400> 4344
gagcaggacc tccattagaa atattaaatt tgattccaaa atctccattg g 51

<210> 4345
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (4346 is other entry)

<221> misc_feature
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<223> Accession number cg43947977

<400> 4345
caagcgaccg atcccgttgg cggcagcgat gcgagcgatg agctggatgg c 51

<210> 4346
<211> 50
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4345 is other entry)

<221> misc_feature
<222> (25)...(26)
<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43947977

<400> 4346
caagcgaccg atcccgttgg cggcacgatg cgagcgatga gctggatggc 50

<210> 4347
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4348 is other entry)

<221> misc_feature
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<223> Accession number cg43948105

<400> 4347
gcctcctact ccccgacaca ggtgcaggag gaactctgga agcactgctc t 51

<210> 4348
<211> 50
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4347 is other entry)

<221> misc_feature
<222> (25)...(26)
<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43948105

<400> 4348
gcctcctact ccccgacaca ggtgcggagg aactctggaa gcactgctct 50

<210> 4349
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4350 is other entry)

<221> misc_feature

<222> (0)...(0)
<223> Accession number cg43948257

<400> 4349
gcggctcatg cctataatcc cagcattttg ggaggccaaa gcaggaggat c 51

<210> 4350
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4349 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43948257

<400> 4350
gcggctcatg cctataatcc cagcactttg ggaggccaaa gcaggaggat c 51

<210> 4351
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4352 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43948257

<400> 4351
aaattagcca ggtgtggtgg cctgtgcctg tagtcccagc tatttgggag g 51

<210> 4352
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4351 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43948257

<400> 4352
aaattagcca ggtgtggtgg cctgtacctg tagtcccagc tatttgggag g 51

<210> 4353
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4354 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43948280

<400> 4353
acaaatgtcc atcacagagt tttccttttt ttttttttga gacagagtct t 51

<210> 4354
<211> 50
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4353 is other entry)

<221> misc_feature
<222> (25)...(26)
<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43948280

<400> 4354
acaaatgtcc atcacagagt tttccttttt tttttttgag acagagtctt 50

<210> 4355
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4356 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43948730

<400> 4355
gctcacggtt ggaggccacg cgctcgtaca gctctgccgc tttggggaac a 51

<210> 4356
<211> 51

<212> DNA
<213> Homo sapiens
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<223> 2 of 2 allelic variants (4355 is other entry)

<221> misc_feature
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<223> Accession number cg43948730

<400> 4356
gctcacggtt ggaggccacg cgctcataca gctctgccgc ttggggaac a 51

<210> 4357
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4358 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43948766

<400> 4357
agtgggtgaca ctgcttgtgt tagtacgccg gggtgcgttg cgtgcggtct c 51

<210> 4358
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4357 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43948766

<400> 4358
agtgggtgaca ctgcttgtgt tagtatgccg gggtgcgttg cgtgcggtct c 51

<210> 4359
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (4360 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43948766

<400> 4359

tctagctgca gctgcatctg cagttgttgt aactgagaag gggaagggcc a

51

<210> 4360

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (4359 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43948766

<400> 4360

tctagctgca gctgcatctg cagttattgt aactgagaag gggaagggcc a

51

<210> 4361

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (4362 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43949166

<400> 4361

cttcaaattgg tgcagttggt taaatgtgaa gaagatgctg cccaggcagt a

51

<210> 4362

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (4361 is other entry)

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<222> (0)...(0)

<223> Accession number cg43949166

<400> 4362
cttcaaattgg tgcagttggt taaatatgaa gaagatgctg cccaggcagt a 51

<210> 4363
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4364 is other entry)

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<223> Accession number cg43949166

<400> 4363
aaaacaaaga tgtgaagaca tggtagatgt gccaagggtta aagatgcttc a 51

<210> 4364
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4363 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43949166

<400> 4364
aaaacaaaga tgtgaagaca tggtaaatgt gccaagggtta aagatgcttc a 51

<210> 4365
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4366 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43949166

<400> 4365
acatggtaga tgtgccaagg ttaaagatgc ttcaaattggt gcagttgttt a 51

<210> 4366
<211> 51
<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (4365 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43949166

<400> 4366

acatggtaga tgtgcaagg ttaaaaatgc ttcaaatggt gcagttgttt a

51

<210> 4367

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (4368 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43949223

<400> 4367

acttgtgaaa gaaggcagca cctgtcagca ccatggacag ctcacaggag t

51

<210> 4368

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (4367 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43949223

<400> 4368

acttgtgaaa gaaggcagca cctgttagca ccatggacag ctcacaggag t

51

<210> 4369

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (4370 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43949223

<400> 4369
ttgtgaaaga aggcagcacc tgtcagcacc atggacagct cacaggagta g 51

<210> 4370
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4369 is other entry)

<221> misc_feature
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<223> Accession number cg43949223

<400> 4370
ttgtgaaaga aggcagcacc tgtcaccacc atggacagct cacaggagta g 51

<210> 4371
<211> 50
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4372 is other entry)

<221> misc_feature
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<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43949223

<400> 4371
gaaagaaggc agcacctgtc agcaccatgg acagctcaca ggagtagttg 50

<210> 4372
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4371 is other entry)

<221> misc_feature

<222> (0)...(0)
<223> Accession number cg43949223

<400> 4372
gaaagaaggc agcacctgtc agcacgcatg gacagctcac aggagtagtt g 51

<210> 4373
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4374 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43949443

<400> 4373
agcggacgtg catcttggtc tcaatatcga tcccctgccca gatctggaag g 51

<210> 4374
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4373 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43949443

<400> 4374
agcggacgtg catcttggtc tcaatgtcga tcccctgccca gatctggaag g 51

<210> 4375
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<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (4376 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43949559

<400> 4375
ttgcccttgg tctcgggggtc gctgtaggcg ctgaggctgc agctatcatg g 51

<210> 4376
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 2 of 2 allelic variants (4375 is other entry)

<221> misc_feature
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<223> Accession number cg43949559

<400> 4376
ttgcccttgg tctcgggggc gctgtcggcg ctgaggctgc agctatcatg g 51

<210> 4377
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (4378 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43949806

<400> 4377
ctcccttacc aaccctgggg ctttatactc cctctccacc aatccctgat g 51

<210> 4378
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 2 of 2 allelic variants (4377 is other entry)

<221> misc_feature
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<223> Accession number cg43949806

<400> 4378
ctcccttacc aaccctgggg ctttagactc cctctccacc aatccctgat g 51

<210> 4379
<211> 51
<212> DNA
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<223> 1 of 2 allelic variants (4380 is other entry)

<221> misc_feature
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<223> Accession number cg43949858

<400> 4379
tcaaaagctg ccgagtccta tgattacacg cgatgggact tgtacacttg a 51

<210> 4380
<211> 51
<212> DNA
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4379 is other entry)

<221> misc_feature
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<223> Accession number cg43949858

<400> 4380
tcaaaagctg ccgagtccta tgattgcacg cgatgggact tgtacacttg a 51

<210> 4381
<211> 51
<212> DNA
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<223> 1 of 2 allelic variants (4382 is other entry)

<221> misc_feature
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<223> Accession number cg43949858

<400> 4381
agctgccgag tcctatgatt acacgcgatg ggacttgtag acttgaagtg a 51

<210> 4382
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 2 of 2 allelic variants (4381 is other entry)

<221> misc_feature
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<223> Accession number cg43949858

<400> 4382

agctgccgag tcctatgatt acacgtgatg ggacttgtag acttgaagtg a

51

<210> 4383

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 1 of 2 allelic variants (4384 is other entry)

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<223> Accession number cg43949935

<400> 4383

aatccatcca caagaaagga gccaaacgtc agtttccaat ggggggaaca t

51

<210> 4384

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 2 of 2 allelic variants (4383 is other entry)

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<223> Accession number cg43949935

<400> 4384

aatccatcca caagaaagga gccaaacgtc agtttccaat ggggggaaca t

51

<210> 4385

<211> 51

<212> DNA

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<223> 1 of 2 allelic variants (4386 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43949954

<400> 4385

cagagaaaac aaggcacttt gggagcatta tggcttactc tactacatgt a

51

<210> 4386

<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4385 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43949954

<400> 4386
cagagaaaac aaggcacttt gggagtatta tggcttactc tactacatgt a 51

<210> 4387
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4388 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43950113

<400> 4387
tggatttttta cctttgcaga cacccaaaaa aaaaataaaa taaatatattt t 51

<210> 4388
<211> 50
<212> DNA
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4387 is other entry)

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<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43950113

<400> 4388
tggatttttta cctttgcaga cacccaaaaa aaaataaaaat aaatatattt 50

<210> 4389
<211> 51
<212> DNA

<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4390 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43950155

<400> 4389
aagagcaaag ccaggaacta gctgagcaga gacccccaaa ccggttggcg g 51

<210> 4390
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (4389 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43950155

<400> 4390
aagagcaaag ccaggaacta gctgaacaga gacccccaaa ccggttggcg g 51

<210> 4391
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4392 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43950281

<400> 4391
caactatatc atcggaggct cggtaatcaa tgagcttatt ggaaatctgg t 51

<210> 4392
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4391 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43950281

<400> 4392
caactatatc atcggaggct cggtagtcaa tgagcttatt ggaaatctgg t 51

<210> 4393
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4394 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43950348

<400> 4393
ccatgcggat catgcgcttg gctacgtcca ggtegccctg gtggttctgg c 51

<210> 4394
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4393 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43950348

<400> 4394
ccatgcggat catgcgcttg gctacatcca ggtegccctg gtggttctgg c 51

<210> 4395
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4396 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43950825

<400> 4395

agtggggacc acaggcaagt ggcaccacga ctaatTTTTT tttatTTTTT g

51

<210> 4396

<211> 51

<212> DNA

<213> Homo sapiens

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<222> (26)...(0)

<223> 2 of 2 allelic variants (4395 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43950825

<400> 4396

agtggggacc acaggcaagt ggcactacga ctaatTTTTT tttatTTTTT g

51

<210> 4397

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (4398 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43950873

<400> 4397

cggtagtggc cccgaatggc tgggcgcgct gatatttatt gcatacaaga c

51

<210> 4398

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (4397 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43950873

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51

<210> 4399

<211> 51

<212> DNA

<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (4400 is other entry)

<221> misc_feature
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<223> Accession number cg43950873

<400> 4399
gcagggtgaag gaggggtgaat cttctaagtg attgacaagg tgaagcaagt c 51

<210> 4400
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (4399 is other entry)

<221> misc_feature
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<223> Accession number cg43950873

<400> 4400
gcagggtgaag gaggggtgaat cttctcagtg attgacaagg tgaagcaagt c 51

<210> 4401
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4402 is other entry)

<221> misc_feature
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<223> Accession number cg43950873

<400> 4401
cttctaagtg attgacaagg tgaagcaagt cacgtgatca taggacagg g 51

<210> 4402
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4401 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43950873

<400> 4402
cttctaagtg attgacaagg tgaagtaagt cacgtgatca taggacaggg g 51

<210> 4403
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4404 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43950873

<400> 4403
gtgattgaca aggtgaagca agtcacgtga tcataggaca gggggccctt c 51

<210> 4404
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (4403 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43950873

<400> 4404
gtgattgaca aggtgaagca agtcacgtga tcataggaca gggggccctt c 51

<210> 4405
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4406 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43950910

<400> 4405
aaaatgcatt ttaacttctg aggggtgggtg tgcaaaatgt tcaccatccc c 51

<210> 4406
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4405 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43950910

<400> 4406
aaaatgcatt ttaacttctg agggttggtg tgcaaaatgt tcaccatccc c 51

<210> 4407
<211> 50
<212> DNA
<213> Homo sapiens

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<222> (26)...(0)
<223> 1 of 2 allelic variants (4408 is other entry)

<221> misc_feature
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<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43950936

<400> 4407
ttcaccatca gtttttcttt tttttctttt tttttttttg cataggcatt 50

<210> 4408
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4407 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43950936

<400> 4408
ttcaccatca gtttttcttt tttttctttt tttttttttt gcataggcat t 51

<210> 4409

<211> 51
<212> DNA
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<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4410 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43950936

<400> 4409
ttttcttttt tttctttttt tttttttgca taggcattac tagggacata a 51

<210> 4410
<211> 50
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4409 is other entry)

<221> misc_feature
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<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43950936

<400> 4410
ttttcttttt tttctttttt tttttttgcat aggcattact agggacataa 50

<210> 4411
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4412 is other entry)

<221> misc_feature
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<223> Accession number cg43950982

<400> 4411
aatgaaaaga ggccccctga aggcctgaa cctggatagg aacaatttgc a 51

<210> 4412
<211> 50
<212> DNA

<213> Homo sapiens

<220>

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<222> (26)...(0)

<223> 2 of 2 allelic variants (4411 is other entry)

<221> misc_feature

<222> (25)...(26)

<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43950982

<400> 4412

aatgaaaaga ggccccctga aggcctgaac ctggatagga acaatttgca

50

<210> 4413

<211> 51

<212> DNA

<213> Homo sapiens

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<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (4414 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43950982

<400> 4413

acgtcacatg gtcaaagtct cctcatttca gccagtctca acacaaaaca c

51

<210> 4414

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (4413 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43950982

<400> 4414

acgtcacatg gtcaaagtct cctcacttca gccagtctca acacaaaaca c

51

<210> 4415

<211> 51

<212> DNA

<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (4416 is other entry)

<221> misc_feature
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<223> Accession number cg43950982

<400> 4415
agttctcaaca caaaacaccc aacagggatg cactcaactt gttggttcca t 51

<210> 4416
<211> 51
<212> DNA
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<220>
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<223> 2 of 2 allelic variants (4415 is other entry)

<221> misc_feature
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<400> 4416
agttctcaaca caaaacaccc aacagagatg cactcaactt gttggttcca t 51

<210> 4417
<211> 51
<212> DNA
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<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4418 is other entry)

<221> misc_feature
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<223> Accession number cg43950982

<400> 4417
gatgcactca acttggttgg tccatgtgga actaggtggc agggcgagag g 51

<210> 4418
<211> 51
<212> DNA
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<220>
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<223> 2 of 2 allelic variants (4417 is other entry)

<221> misc_feature

<222> (0)...(0)
<223> Accession number cg43950982

<400> 4418
gatgcactca acttggttggg tccatttggg actaggtggc agggcgagag g 51

<210> 4419
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (4420 is other entry)

<221> misc_feature
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<223> Accession number cg43950982

<400> 4419
gttggttcca tgtggaacta ggtggcaggg cgagagggaa agtagtagaa g 51

<210> 4420
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<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (4419 is other entry)

<221> misc_feature
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<223> Accession number cg43950982

<400> 4420
gttggttcca tgtggaacta ggtggtaggg cgagagggaa agtagtagaa g 51

<210> 4421
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4422 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43950982

<400> 4421
aggtggcagg gcgagagggg aagtagtaga agggggctat ggtgtgtctg c 51

<210> 4422
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4421 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43950982

<400> 4422
aggtggcagg gcgagaggga aagtattaga agggggctat ggtgtgtctg c 51

<210> 4423
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4424 is other entry)

<221> misc_feature
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<223> Accession number cg43950982

<400> 4423
agggcgagag ggaaagtagt agaagggggc tatggtgtgt ctgcattcag t 51

<210> 4424
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4423 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43950982

<400> 4424
agggcgagag ggaaagtagt agaagcgggc tatggtgtgt ctgcattcag t 51

<210> 4425
<211> 51
<212> DNA
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<220>

<221> misc_feature
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<223> 1 of 2 allelic variants (4426 is other entry)

<221> misc_feature
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<400> 4425
agaggggaaag tagtagaagg gggctatggg gtgtctgcat tcagtcccct c 51

<210> 4426
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4425 is other entry)

<221> misc_feature
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<400> 4426
agaggggaaag tagtagaagg gggctgtggg gtgtctgcat tcagtcccct c 51

<210> 4427
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<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4428 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43950982

<400> 4427
tgtgtctgca ttcagtcccc tcacataaag ccacatggat ctaggggggt a 51

<210> 4428
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<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4427 is other entry)

<221> misc_feature
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<223> Accession number cg43950982

<400> 4428
tgtgtctgca ttcagtcctc tcacacaaag ccacatggat ctaggggggt a 51

<210> 4429
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4430 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43951482

<400> 4429
gctttcaaaa gacattttgt gaaggacatt aatttcacat ttaaaacgtg t 51

<210> 4430
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4429 is other entry)

<221> misc_feature
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<223> Accession number cg43951482

<400> 4430
gctttcaaaa gacattttgt gaaggacatt aatttcacat ttaaaacgtg t 51

<210> 4431
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4432 is other entry)

<221> misc_feature
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<223> Accession number cg43951580

<400> 4431
acacacacac acacacacac acacacacac acaaccttct gtggctcaaa a 51

<210> 4432

<211> 50
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (4431 is other entry)

<221> misc_feature
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<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43951580

<400> 4432
acacacacac acacacacac acacaacaca caaccttctg tggctcaaaa 50

<210> 4433
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 1 of 2 allelic variants (4434 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43951635

<400> 4433
atgcctggct aatttttgta ttttttagtag agacaggggtt tcaccatatt g 51

<210> 4434
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (4433 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43951635

<400> 4434
atgcctggct aatttttgta tttttggttag agacaggggtt tcaccatatt g 51

<210> 4435
<211> 51
<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

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<223> 1 of 2 allelic variants (4436 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43951719

<400> 4435

gtggcaggat aaaaggatat ttgtgaagta atcttagggt tggataaaaa g

51

<210> 4436

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

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<223> 2 of 2 allelic variants (4435 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43951719

<400> 4436

gtggcaggat aaaaggatat ttgtggagta atcttagggt tggataaaaa g

51

<210> 4437

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

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<223> 1 of 2 allelic variants (4438 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43951812

<400> 4437

gcaggtacag gtgccagttt gtgacggatg aaagcacga cagcccacgc g

51

<210> 4438

<211> 51

<212> DNA

<213> Homo sapiens

<220>

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<222> (26)...(0)

<223> 2 of 2 allelic variants (4437 is other entry)

<221> misc_feature
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<223> Accession number cg43951812

<400> 4438
gcaggtacag gtgccagttt gtgacagatg aaagcaccga cagcccacgc g 51

<210> 4439
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4440 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43951812

<400> 4439
accccagaag caaaatgctc catgcaacag ccaggcattc aggctacaag c 51

<210> 4440
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (4439 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43951812

<400> 4440
accccagaag caaaatgctc catgcgacag ccaggcattc aggctacaag c 51

<210> 4441
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4442 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43951883

<400> 4441

atttacagac ttgagtgtgt gtgtgtgttt ccaaccacag tcattcatac t

51

<210> 4442

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 2 of 2 allelic variants (4441 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43951883

<400> 4442

atttacagac ttgagtgtgt gtgtgggttt ccaaccacag tcattcatac t

51

<210> 4443

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 1 of 2 allelic variants (4444 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43952028

<400> 4443

ccctggtgtg tagctagcaa gcaataactg actactcgtc acctacagtt g

51

<210> 4444

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (4443 is other entry)

<221> misc_feature

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<223> Accession number cg43952028

<400> 4444

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51

<210> 4445

<211> 51

<212> DNA

<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (4446 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43952230

<400> 4445
actgcccaatt gctcattttg tctgatatta acagattatg catttcctca g 51

<210> 4446
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4445 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43952230

<400> 4446
actgcccaatt gctcattttg tctgacatta acagattatg catttcctca g 51

<210> 4447
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4448 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43952230

<400> 4447
ctgttcattg tccacatgta ttaaacaata aacatgtcaa ttacttggtg c 51

<210> 4448
<211> 50
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (4447 is other entry)

<221> misc_feature
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<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43952230

<400> 4448
ctgttcacatgga tccacatgta ttaaaaaaaaa acatgtcaat tacttggtgc 50

<210> 4449
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4450 is other entry)

<221> misc_feature
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<223> Accession number cg43952230

<400> 4449
tcattggtcca catgtattaa acaaaaaaaca tgtcaattac ttggtgcaaa c 51

<210> 4450
<211> 50
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (4449 is other entry)

<221> misc_feature
<222> (25)...(26)
<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43952230

<400> 4450
tcattggtcca catgtattaa acaaaaacat gtcaattact ttggtgcaaac 50

<210> 4451
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (4452 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43952230

<400> 4451

catggtccac atgtattaaa caaaaaacat gtcaattact tggtgcaaac a

51

<210> 4452

<211> 50

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (4451 is other entry)

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<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43952230

<400> 4452

catggtccac atgtattaaa caaaaacatg tcaattactt ggtgcaaaca

50

<210> 4453

<211> 51

<212> DNA

<213> Homo sapiens

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<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (4454 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43952230

<400> 4453

atggtccaca tgtattaaac aaaaaacatg tcaattactt ggtgcaaaca c

51

<210> 4454

<211> 50

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (4453 is other entry)

<221> misc_feature
<222> (25)...(26)
<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43952230

<400> 4454
atggtccaca tgtattaaac aaaaacatgt caattacttg gtgcaaacac 50

<210> 4455
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4456 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43952399

<400> 4455
agggcactct gtccttgca ataacatcct actgaacagt aagtaccatg g 51

<210> 4456
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4455 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43952399

<400> 4456
agggcactct gtccttgca ataacgtcct actgaacagt aagtaccatg g 51

<210> 4457
<211> 50
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4458 is other entry)

<221> misc_feature
<222> (25)...(26)

<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43953987

<400> 4457

tactaaaagg gaggggggca tgctttccaaa tggggatcta cgtcttcctc

50

<210> 4458

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (4457 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43953987

<400> 4458

tactaaaagg gaggggggca tgctttccaa atggggatct acgtcttcct c

51

<210> 4459

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (4460 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43955058

<400> 4459

gctgcaggga gctggaagtg gtcagtatat atccgagccg ctgtccgaaa a

51

<210> 4460

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (4459 is other entry)

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<222> (0)...(0)

<223> Accession number cg43955058

<400> 4460
gctgcaggga gctggaagtg gtcagcatat atccgagccg ctgtccgaaa a 51

<210> 4461
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4462 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43955367

<400> 4461
aggagttcaa gaccagcctg accaacaatgg tgaaacccca tgtctactaa a 51

<210> 4462
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4461 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43955367

<400> 4462
aggagttcaa gaccagcctg accaatatgg tgaaacccca tgtctactaa a 51

<210> 4463
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4464 is other entry)

<221> misc_feature
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<223> Accession number cg43955367

<400> 4463
ggagttcaag accagcctga ccaacatggg gaaaccccat gtctactaaa a 51

<210> 4464
<211> 51
<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (4463 is other entry)

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<222> (0)...(0)

<223> Accession number cg43955367

<400> 4464

ggagttcaag accagcctga ccaacgtggt gaaaccccat gtctactaaa a

51

<210> 4465

<211> 51

<212> DNA

<213> Homo sapiens

<220>

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<222> (26)...(0)

<223> 1 of 2 allelic variants (4466 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43955367

<400> 4465

ctactaaaaa tacaaaaaaa tgagccgggc atggtggcgc gtgcctgtaa t

51

<210> 4466

<211> 51

<212> DNA

<213> Homo sapiens

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<222> (26)...(0)

<223> 2 of 2 allelic variants (4465 is other entry)

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<400> 4466

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51

<210> 4467

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (4468 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43955367

<400> 4467
aaaaatacaa aaaaatgagc cgggcatggt ggcgcgtgcc tgtaatccca g 51

<210> 4468
<211> 51
<212> DNA
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<220>
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<223> 2 of 2 allelic variants (4467 is other entry)

<221> misc_feature
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<223> Accession number cg43955367

<400> 4468
aaaaatacaa aaaaatgagc cgggcgtggt ggcgcgtgcc tgtaatccca g 51

<210> 4469
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4470 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43955553

<400> 4469
gaaatctaag gcaagtaaag gcacaatgaa tgtggagttg aatggaacca a 51

<210> 4470
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4469 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43955553

<400> 4470

gaaatctaata gcaagtaaag gcacagtgaatgtgttggaacca a

51

<210> 4471

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (4472 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43955835

<400> 4471

gtcacagacc actcgccatt gagaaaagac agagtttggg cttattagag t

51

<210> 4472

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (4471 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43955835

<400> 4472

gtcacagacc actcgccatt gagaagagac agagtttggg cttattagag t

51

<210> 4473

<211> 51

<212> DNA

<213> Homo sapiens

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<222> (26)...(0)

<223> 1 of 2 allelic variants (4474 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43955871

<400> 4473

ttcccagtgt gtggtcttct ttgttgatgg tgaataaaca ggcattgggt a

51

<210> 4474

<211> 51

<212> DNA

<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4473 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43955871

<400> 4474
ttcccagtggt gtggtcttct ttgttcatgg tgaataaaca ggcattgggt a 51

<210> 4475
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4476 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43955877

<400> 4475
tcaaagtcct attgtaatat tatTTtaagg gtcttaggag gccctcaga g 51

<210> 4476
<211> 50
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4475 is other entry)

<221> misc_feature
<222> (25)...(26)
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<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43955877

<400> 4476
tcaaagtcct attgtaatat tatTTaaggg tcttaggagg cccctcagag 50

<210> 4477
<211> 51
<212> DNA
<213> Homo sapiens

<220>

<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4478 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43955877

<400> 4477
ggtcagggct agagtatgag aagtcctaag ggtttttgta ttttgttttt t 51

<210> 4478
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4477 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43955877

<400> 4478
ggtcagggct agagtatgag aagtcataag ggtttttgta ttttgttttt t 51

<210> 4479
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4480 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43955877

<400> 4479
aagggttttt gtattttggt tttttttcct ataaaccctg aggttgaaag c . 51

<210> 4480
<211> 50
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4479 is other entry)

<221> misc_feature
<222> (25)...(26)

<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43955877

<400> 4480

aagggttttt gtattttggt ttttttccta taaacctga ggttgaaagc

50

<210> 4481

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (4482 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43955877

<400> 4481

gtattttggt ttttttcct ataaacctg aggttgaaag ctctggatag c

51

<210> 4482

<211> 50

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (4481 is other entry)

<221> misc_feature

<222> (25)...(26)

<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43955877

<400> 4482

gtattttggt ttttttcct ataaacctga ggttgaaagc tctggatagc

50

<210> 4483

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (4484 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43956161

<400> 4483
agaggctggt attccaggga aagagcgggc aggtgtcctg aggagtacat t 51

<210> 4484
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (4483 is other entry)

<221> misc_feature
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<223> Accession number cg43956161

<400> 4484
agaggctggt attccaggga aagagtgggc aggtgtcctg aggagtacat t 51

<210> 4485
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4486 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43956227

<400> 4485
tttctgaaat tcacaaagtt aaacgtgatg tgctcatcag aaacaatttc t 51

<210> 4486
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4485 is other entry)

<221> misc_feature
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<223> Accession number cg43956227

<400> 4486
tttctgaaat tcacaaagtt aaacgcgatg tgctcatcag aaacaatttc t 51

<210> 4487
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4488 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43956347

<400> 4487
tcaacaatta gcagcttaag atctatcaac tacagtgtta acgttcacac g 51

<210> 4488
<211> 51
<212> DNA
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<223> 2 of 2 allelic variants (4487 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43956347

<400> 4488
tcaacaatta gcagcttaag atctagcaac tacagtgtta acgttcacac g 51

<210> 4489
<211> 50
<212> DNA
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<223> 1 of 2 allelic variants (4490 is other entry)

<221> misc_feature
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<221> misc_feature
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<400> 4489
tcaactacag tggttaacgtt cacacgttca caagtgtcat ttctttacgt 50

<210> 4490

<211> 51
<212> DNA
<213> Homo sapiens

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<223> 2 of 2 allelic variants (4489 is other entry)

<221> misc_feature
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<223> Accession number cg43956347

<400> 4490
tcaactacag tggttaacgtt cacacagttc acaagtgta tttctttacg t 51

<210> 4491
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 1 of 2 allelic variants (4492 is other entry)

<221> misc_feature
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<223> Accession number cg43956347

<400> 4491
tacagtgtta acgttcacac gttcacaagt gtcatttctt tacgtttcaa t 51

<210> 4492
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 2 of 2 allelic variants (4491 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43956347

<400> 4492
tacagtgtta acgttcacac gttcataagt gtcatttctt tacgtttcaa t 51

<210> 4493
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 1 of 2 allelic variants (4494 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43956347

<400> 4493
cagtgttaac gttcacacgt tcacaagtgt catttcttta cgtttcaatt c 51

<210> 4494
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 2 of 2 allelic variants (4493 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43956347

<400> 4494
cagtgttaac gttcacacgt tcacaggtgt catttcttta cgtttcaatt c 51

<210> 4495
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 1 of 2 allelic variants (4496 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43956467

<400> 4495
cctcactatc gagatacttg tgggtggcgt agcccatcag ggcaccaata t 51

<210> 4496
<211> 50
<212> DNA
<213> Homo sapiens

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<223> 2 of 2 allelic variants (4495 is other entry)

<221> misc_feature
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<221> misc_feature
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<223> Accession number cg43956467

<400> 4496
cctcactatc gagatacttg tgggtgcgta gcccatcagg gcaccaatat 50

<210> 4497
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 1 of 2 allelic variants (4498 is other entry)

<221> misc_feature
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<223> Accession number cg43956467

<400> 4497
ctcactatcg agatacttgt ggggtggcgta gcccatcagg gcaccaatat t 51

<210> 4498
<211> 50
<212> DNA
<213> Homo sapiens

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<223> 2 of 2 allelic variants (4497 is other entry)

<221> misc_feature
<222> (25)...(26)
<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
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<223> Accession number cg43956467

<400> 4498
ctcactatcg agatacttgt ggggtgcgtag cccatcaggg caccaatatt 50

<210> 4499
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (4500 is other entry)

<221> misc_feature

<222> (0)...(0)
<223> Accession number cg43956696

<400> 4499
cactcctact tctctcaaatt tagcatttta acatctttcca ataacacacc a 51

<210> 4500
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4499 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43956696

<400> 4500
cactcctact tctctcaaatt tagcatttta acatctttcca ataacacacc a 51

<210> 4501
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4502 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43956870

<400> 4501
aggatggctg ggttgaggtc ttgctcaag ttctgctaac attgatgaca g 51

<210> 4502
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4501 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43956870

<400> 4502
aggatggctg ggttgaggtc ttgcccgaag ttctgctaac attgatgaca g 51

<210> 4503
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4504 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43956870

<400> 4503
aattgttctc atccattaac gctgctgtgt ctttctggct ctttgcaaaa g 51

<210> 4504
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4503 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43956870

<400> 4504
aattgttctc atccattaac gctgccgtgt ctttctggct ctttgcaaaa g 51

<210> 4505
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4506 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43957151

<400> 4505
aagtccaaca tctcgcccaa cttcaacttc atggggcagc tgctggactt t 51

<210> 4506
<211> 51
<212> DNA
<213> Homo sapiens

<220>

<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4505 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43957151

<400> 4506
aagtccaaca tctcgcccaa cttcagcttc atggggcagc tgctggactt t 51

<210> 4507
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4508 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43957194

<400> 4507
acaaggcttc ttgtctcagg tctgcagtgt gtcacatgcc agactcctca g 51

<210> 4508
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4507 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43957194

<400> 4508
acaaggcttc ttgtctcagg tctgcggtgt gtcacatgcc agactcctca g 51

<210> 4509
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4510 is other entry)

<221> misc_feature
<222> (0)...(0)

<223> Accession number cg43957358

<400> 4509

tgggctccac caccgtggcc gccggcggga ccagcacagg cggcgttttc t

51

<210> 4510

<211> 50

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (4509 is other entry)

<221> misc_feature

<222> (25)...(26)

<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43957358

<400> 4510

tgggctccac caccgtggcc gccggggggac cagcacaggc ggcgttttct

50

<210> 4511

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (4512 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43957502

<400> 4511

agctggaggg aacgccagag gtgtcctgcc gggctctgga gctcttcgac t

51

<210> 4512

<211> 50

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

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<223> 2 of 2 allelic variants (4511 is other entry)

<221> misc_feature

<222> (25)...(26)

<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43957502

<400> 4512
agctggaggg aacgccagag gtgtctgccg ggctctggag ctcttcgact 50

<210> 4513
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4514 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43957567

<400> 4513
caaatgacta caatgttaaa atagacaaaa actgctatac aagagcctct t 51

<210> 4514
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (4513 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43957567

<400> 4514
caaatgacta caatgttaaa atagataaaa actgctatac aagagcctct t 51

<210> 4515
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4516 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43957735

<400> 4515
aaaccctttt gagcttgagt aaccactgag ctgccctttg ttacctttat g 51

<210> 4516
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4515 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43957735

<400> 4516
aaaccctttt gagcttgagt aaccattgag ctgccctttg ttacctttat g 51

<210> 4517
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4518 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43957996

<400> 4517
aatatcttgg aaatccacat cattcacagc tagaacttgg tccccttctt g 51

<210> 4518
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4517 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43957996

<400> 4518
aatatcttgg aaatccacat cattgcgagc tagaacttgg tccccttctt g 51

<210> 4519
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4520 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43958045

<400> 4519
ctcagctagt ccagaaattg ctgcatttcc catattactt agttctttat t 51

<210> 4520
<211> 50
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4519 is other entry)

<221> misc_feature
<222> (25)...(26)
<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43958045

<400> 4520
ctcagctagt ccagaaattg ctgcattccc atattactta gttctttatt 50

<210> 4521
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (4522 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43958093

<400> 4521
ggtcttgctc tttatgagag ggcaatgtgt ttttaattgt gttaattaga a 51

<210> 4522
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature

<222> (26)...(0)
<223> 2 of 2 allelic variants (4521 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43958093

<400> 4522
ggctcttgctc tttatgagag ggcaacgtgt ttttaattgt gttaattaga a 51

<210> 4523
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4524 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43958290

<400> 4523
actaaggagg ctgaggtagg agaatgactt gaacctggga ggcagaggtt g 51

<210> 4524
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4523 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43958290

<400> 4524
actaaggagg ctgaggtagg agaatcactt gaacctggga ggcagaggtt g 51

<210> 4525
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4526 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43958290

<400> 4525
ggagaatgac ttgaacctgg gaggcagagg ttgtagtgag ctgagatctt g 51

<210> 4526
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4525 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43958290

<400> 4526
ggagaatgac ttgaacctgg gaggcggagg ttgtagtgag ctgagatctt g 51

<210> 4527
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4528 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43958316

<400> 4527
ggtcgggga gcaatggcat cctccttgat atcattggct ggctcctcca g 51

<210> 4528
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4527 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43958316

<400> 4528
ggtcgggga gcaatggcat cctccgtgat atcattggct ggctcctcca g 51

<210> 4529
<211> 51

<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4530 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43958316

<400> 4529
attggctggc tctccaggg ccttgcccg gcggtgtac aagaggacca g

51

<210> 4530
<211> 50
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (4529 is other entry)

<221> misc_feature
<222> (25)...(26)
<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43958316

<400> 4530
attggctggc tctccaggg ccttgcccg cggtgtaca agaggaccag

50

<210> 4531
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4532 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43958676

<400> 4531
attctttaa ttccaatc ctcttatctt cctcctctc ctctgccga g

51

<210> 4532
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4531 is other entry)

<221> misc_feature
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<223> Accession number cg43958676

<400> 4532
attctttaa ttctccaatc ctcttggttt cctctctctc ctctgccgca g 51

<210> 4533
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4534 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43958676

<400> 4533
ctttaaattc tccaatcctc ttattttcct cctcttcctc tgccgcagtc a 51

<210> 4534
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4533 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43958676

<400> 4534
ctttaaattc tccaatcctc ttatttctct cctcttcctc tgccgcagtc a 51

<210> 4535
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4536 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43958676

<400> 4535
taaattctcc aatcctctta ttttctctct ctctctctgc cgcagtcact c 51

<210> 4536
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4535 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43958676

<400> 4536
taaattctcc aatcctctta ttttctctct ctctctctgc cgcagtcact c 51

<210> 4537
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4538 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43958676

<400> 4537
cgcagtcact cccactggc cctcaatttc acgctgtttc tttttctctt c 51

<210> 4538
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4537 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43958676

<400> 4538
cgcagtcact cccactggc cctcagtttc acgctgtttc tttttctctt c 51

<210> 4539
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4540 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43958676

<400> 4539
ggccctcaat ttcacgctgt ttctttttct cttcaaactg caatcgctc t 51

<210> 4540
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4539 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43958676

<400> 4540
ggccctcaat ttcacgctgt ttctttttct cttcaaactg caatcgctc t 51

<210> 4541
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4542 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43958676

<400> 4541
cacgctgttt cttttttctt tcaaactgca atcgctctg caactcctcc a 51

<210> 4542
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4541 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43958676 4

<400> 4542
cacgctgttt ctttttctct tcaaattgca atcgctctg caactcctcc a 51

<210> 4543
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4544 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43958676

<400> 4543
tcttcaaact gcaatgcct ctgcaactcc tccagagcag ctctgtacat t 51

<210> 4544
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<212> DNA
<213> Homo sapiens

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<223> 2 of 2 allelic variants (4543 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43958676

<400> 4544
tcttcaaact gcaatgcct ctgcagctcc tccagagcag ctctgtacat t 51

<210> 4545
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (4546 is other entry)

<221> misc_feature

<222> (0)...(0)
<223> Accession number cg43958676

<400> 4545
tcctccagag cagctctgta cattgcgtct tgggcattct gaagttcaat g 51

<210> 4546
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4545 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43958676

<400> 4546
tcctccagag cagctctgta cattgtgtct tgggcattct gaagttcaat g 51

<210> 4547
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4548 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43958676

<400> 4547
cagctctgta cattgcgtct tgggcattct gaagttcaat gatttgatca a 51

<210> 4548
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4547 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43958676

<400> 4548
cagctctgta cattgcgtct tgggcgttct gaagttcaat gatttgatca a 51

<210> 4549
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4550 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43958676

<400> 4549
tgggtgtccag gaacgcctcg tgtgcagcaa taatgtgatc caaatcctgg g 51

<210> 4550
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4549 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43958676

<400> 4550
tgggtgtccag gaacgcctcg tgtgccgcaa taatgtgatc caaatcctgg g 51

<210> 4551
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4552 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43958676

<400> 4551
tgtccaggaa cgcctcgtgt gcagcaataa tgtgatccaa atcctggggc t 51

<210> 4552
<211> 51
<212> DNA
<213> Homo sapiens

<220>

<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4551 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43958676

<400> 4552
tgtccaggaa cgcctcgtgt gcagcgataa tgtgatccaa atcctgggcc t 51

<210> 4553
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4554 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43958676

<400> 4553
caataatgtg atccaaatcc tgggcctgct ggactctgtt ccaaagctca t 51

<210> 4554
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4553 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43958676

<400> 4554
caataatgtg atccaaatcc tgggcttgct ggactctgtt ccaaagctca t 51

<210> 4555
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4556 is other entry)

<221> misc_feature
<222> (0)...(0)

<223> Accession number cg43958676

<400> 4555

cctgggacctg ctggactctg ttccaaagct catccaaga acactcaagc a

51

<210> 4556

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (4555 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43958676

<400> 4556

cctgggacctg ctggactctg ttccagagct catccaaga acactcaagc a

51

<210> 4557

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (4558 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43958736

<400> 4557

atacaataaa tatatcaatt gtttacattc ccaaattttg aaaatactgg g

51

<210> 4558

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (4557 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43958736

<400> 4558

atacaataaa tatatcaatt gtttatattc ccaaattttg aaaatactgg g

51

<210> 4559

<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4560 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43958770

<400> 4559
aagaaaatat ttacaaaata caagggttttt tttttccatt ttttgttttt g 51

<210> 4560
<211> 50
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4559 is other entry)

<221> misc_feature
<222> (25)...(26)
<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43958770

<400> 4560
aagaaaatat ttacaaaata caagggttttt ttttccattt tttgtttttg 50

<210> 4561
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4562 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43958770

<400> 4561
atttttatag gattacaaaa tggccaaaaa aaaagagtct tctccccct c 51

<210> 4562
<211> 50
<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (4561 is other entry)

<221> misc_feature

<222> (25)...(26)

<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43958770

<400> 4562

atttttatac gattacaaaaa tggccaaaaa aaagagtctt ctccccctc

50

<210> 4563

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (4564 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43958939

<400> 4563

gtgtgtataa gtacatcctt tgggggtttt ttttctctt tttttaacc a

51

<210> 4564

<211> 50

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (4563 is other entry)

<221> misc_feature

<222> (25)...(26)

<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43958939

<400> 4564

gtgtgtataa gtacatcctt tgggggtttt ttttctctt tttttaacca

50

<210> 4565

<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4566 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43958939

<400> 4565
agtacatcct ttgggggtttt ttttttctct tttttttaac caaagttgct g 51

<210> 4566
<211> 50
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4565 is other entry)

<221> misc_feature
<222> (25)...(26)
<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43958939

<400> 4566
agtacatcct ttgggggtttt ttttttctct ttttttaacc aaagttgctg 50

<210> 4567
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4568 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43959547

<400> 4567
tgggagtgtc ttttagcatg ctaatgcatt ataattagca tataatgaac t 51

<210> 4568
<211> 51
<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (4567 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43959547

<400> 4568

tgggagtgtc ttttagcatg ctaatacatt ataattagca tataatgaac t

51

<210> 4569

<211> 51

<212> DNA

<213> Homo sapiens

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<222> (26)...(0)

<223> 1 of 2 allelic variants (4570 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43959551

<400> 4569

agttttcaaa gttaatttgg cgggtagggg cttgggaagt ggggagtgt g

51

<210> 4570

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (4569 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43959551

<400> 4570

agttttcaaa gttaatttgg cgggtggggg cttgggaagt ggggagtgt g

51

<210> 4571

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (4572 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43959551

<400> 4571
tgatcgggtca ggttggagat ggattcatag ggggttgaag tgagtttttc a 51

<210> 4572
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4571 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43959551

<400> 4572
tgatcgggtca ggttggagat ggatttatag ggggttgaag tgagtttttc a 51

<210> 4573
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4574 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43959563

<400> 4573
agtcactgtg ccctgagcgg agagcctcag cgtagccgcg gcagccatcg a 51

<210> 4574
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4573 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43959563

<400> 4574

agtcactgtg ccctgagcgg agagcttcag cgtagccgcg gcagccatcg a 51

<210> 4575

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (4576 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43959715

<400> 4575

gaccattcca attcatcttc agctgccaag tgtatttagt ccctgaacct g 51

<210> 4576

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (4575 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43959715

<400> 4576

gaccattcca attcatcttc agctgacaag tgtatttagt ccctgaacct g 51

<210> 4577

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (4578 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43959715

<400> 4577

tccaattcat cttcagctgc caagtgtatt tagtccctga acctggatcc a 51

<210> 4578

<211> 51

<212> DNA

<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4577 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43959715

<400> 4578
tccaattcat cttcagctgc caagtttatt tagtcctga acctggatcc a 51

<210> 4579
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4580 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43959715

<400> 4579
catcttcagc tgccaagtgt atttagtccc tgaacctgga tccaaggcat c 51

<210> 4580
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4579 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43959715

<400> 4580
catcttcagc tgccaagtgt atttaatccc tgaacctgga tccaaggcat c 51

<210> 4581
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4582 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43959715

<400> 4581
ctggatccaa ggcattctccc tgtaggaaac atcagaccgg ggcagagatt g 51

<210> 4582
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4581 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43959715

<400> 4582
ctggatccaa ggcattctccc tgtagaaaac atcagaccgg ggcagagatt g 51

<210> 4583
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4584 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43960167

<400> 4583
agacaaatgc ctaggcagat aggggcaggt caacagtga accccacctc c 51

<210> 4584
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4583 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43960167

<400> 4584
agacaaatgc ctaggcagat aggggtaggt caacagtga accccacctc c 51

<210> 4585
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4586 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43960464

<400> 4585
ggcaagattc cgaatgccag gcccctcaag tgtgcaacag ggcacagggt g 51

<210> 4586
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4585 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43960464

<400> 4586
ggcaagattc cgaatgccag gccccccaag tgtgcaacag ggcacagggt g 51

<210> 4587
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4588 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43960464

<400> 4587
tccgaatgcc aggccctca agtgtgcaac agggcacagg gtgacctcat g 51

<210> 4588
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4587 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43960464

<400> 4588
tccgaatgcc aggccctca agtgtacaac agggcacagg gtgacctcat g 51

<210> 4589
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4590 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43960464

<400> 4589
gtgcaacagg gcacaggggtg acctcatgtg ggcaggtggg tgctgttctg t 51

<210> 4590
<211> 50
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4589 is other entry)

<221> misc_feature
<222> (25)...(26)
<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43960464

<400> 4590
gtgcaacagg gcacaggggtg acctctgtgg gcaggtgggt gctgttctgt 50

<210> 4591
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature

<222> (26)...(0)
<223> 1 of 2 allelic variants (4592 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43960807

<400> 4591
tgctcgattc cacgtgtgct cggggccccc agaagaactc atactccacc g 51

<210> 4592
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4591 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43960807

<400> 4592
tgctcgattc cacgtgtgct cgggggtcccc agaagaactc atactccacc g 51

<210> 4593
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4594 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43960917

<400> 4593
gcccgagca atggaagtct catccccatc ctgagcggcc tcttttctag g 51

<210> 4594
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4593 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43960917

<400> 4594
gccccggagca atggaagtct catccgcatc ctgagcggcc tcttttctag g 51

<210> 4595
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4596 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43960917

<400> 4595
aagtctcatc cccatcctga gcggcctctt ttctaggatc gagaggacca c 51

<210> 4596
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4595 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43960917

<400> 4596
aagtctcatc cccatcctga gcggcctctt ttctaggatc gagaggacca c 51

<210> 4597
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4598 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43960917

<400> 4597
tttctaggat cgagaggacc aactgcagc ccaggacaaa agcccacggt a 51

<210> 4598
<211> 51

<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4597 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43960917

<400> 4598
tttctaggat cgagaggacc acactccagc ccaggacaaa agcccacggt a

51

<210> 4599
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4600 is other

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43960917

<400> 4599
gaccacactg cagcccagga caaaagccca cggtagcaca ttgtccggca g

51

<210> 4600
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4599 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43960917

<400> 4600
gaccacactg cagcccagga caaaatccca cggtagcaca ttgtccggca g

51

<210> 4601
<211> 51
<212> DNA
<213> Homo sapiens

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<221> misc_feature
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<223> 1 of 2 allelic variants (4602 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43960917

<400> 4601

accacactgc agcccaggac aaaagccac ggtagcacat tgtccggcag g

51

<210> 4602

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (4601 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43960917

<400> 4602

accacactgc agcccaggac aaaaggccac ggtagcacat tgtccggcag g

51

<210> 4603

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (4604 is other entry)

<221> misc_feature

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<223> Accession number cg43960917

<400> 4603

actgcagccc aggacaaaag cccacggtag cacattgtcc ggcaggagag g

51

<210> 4604

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 2 of 2 allelic variants (4603 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43960917

<400> 4604
actgcagccc aggacaaaag cccacagtag cacattgtcc ggcaggagag g 51

<210> 4605
<211> 51
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<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4606 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43960917

<400> 4605
gccaggaca aaagcccacg gtagcacatt gtccggcagg agaggagcag a 51

<210> 4606
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4605 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43960917

<400> 4606
gccaggaca aaagcccacg gtagctcatt gtccggcagg agaggagcag a 51

<210> 4607
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4608 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43960917

<400> 4607
aaaagccac ggtagcacat tgtccggcag gagaggagca gaccacgtc c 51

<210> 4608
<211> 51
<212> DNA

<213> Homo sapiens

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<222> (26)...(0)
<223> 2 of 2 allelic variants (4607 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43960917

<400> 4608
aaaagcccac ggtagcacat tgtccagcag gagaggagca gacccacgtc c 51

<210> 4609
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (4610 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43960917

<400> 4609
tcgggcagga gaggagcaga cccacgtcca agaagatggt ttacctttg c 51

<210> 4610
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (4609 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43960917

<400> 4610
tcgggcagga gaggagcaga cccacctcca agaagatggt ttacctttg c 51

<210> 4611
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (4612 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43960917

<400> 4611
cagacccacg tccaagaaga tggttttacc tttgcacgcc tcttctctga g 51

<210> 4612
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (4611 is other entry)

<221> misc_feature
<222> (0)...(0)
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<400> 4612
cagacccacg tccaagaaga tggttgtacc tttgcacgcc tcttctctga g . 51

<210> 4613
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (4614 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43960917

<400> 4613
ccacgtccaa gaagatgggtt ttacctttgc acgcctcttc tctgagaaat g 51

<210> 4614
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 2 of 2 allelic variants (4613 is other entry)

<221> misc_feature
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<223> Accession number cg43960917

<400> 4614

ccacgtccaa gaagatgggt ttacccttgc acgcctcttc tctgagaaat g 51

<210> 4615

<211> 51

<212> DNA

<213> Homo sapiens

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<222> (26)...(0)

<223> 1 of 2 allelic variants (4616 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43961283

<400> 4615

ccatgcctcc tagcaagatg ctgaggctac agtaggtctg gcctcaagct g 51

<210> 4616

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 2 of 2 allelic variants (4615 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43961283

<400> 4616

ccatgcctcc tagcaagatg ctgagcctac agtaggtctg gcctcaagct g 51

<210> 4617

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 1 of 2 allelic variants (4618 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43961690

<400> 4617

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<210> 4618

<211> 51

<212> DNA

<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (4617 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43961690

<400> 4618
ctgaggcagg agaattgctt gaacctggga ggcagagggtt gcagtgagcc g 51

<210> 4619
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4620 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43961969

<400> 4619
ctattagtaa acaaggccta cgtttttttc tctaaaattt agaatcttaa a 51

<210> 4620
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4619 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43961969

<400> 4620
ctattagtaa acaaggccta cgtttggttc tctaaaattt agaatcttaa a 51

<210> 4621
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4622 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43961990

<400> 4621
cttgccctcat ccctgtcttt ggcaagtgcg cgggtggtgt ggaggaaagg a 51

<210> 4622
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4621 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43961990

<400> 4622
cttgccctcat ccctgtcttt ggcaactgcg cgggtggtgt ggaggaaagg a 51

<210> 4623
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4624 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43962112

<400> 4623
ctccaggag gctcagatca tggttcaggg gtgccaggca ccattcctac t 51

<210> 4624
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4623 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43962112

<400> 4624
ctccaggag gctcagatca tggtttaggg gtgccaggca ccattcctac t 51

<210> 4625
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4626 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43962250

<400> 4625
tcatgccacc tgcgagacgg gctcctctctg tccccactgt gtccccgggt c 51

<210> 4626
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4625 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43962250

<400> 4626
tcatgccacc tgcgagacgg gctccccctg tccccactgt gtccccgggt c 51

<210> 4627
<211> 51
<212> DNA
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<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4628 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43962322

<400> 4627
gtgtgggaag ggccagaata agcaacaaag ccaattagat gtgggttctg g 51

<210> 4628
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4627 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43962322

<400> 4628
gtgtgggaag ggccagaata agcaataaag ccaattagat gtgggttctg g 51

<210> 4629
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4630 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43962689

<400> 4629
tactgagggt ctatgcatgg taattggcag aaaacatgat agaactaaac a 51

<210> 4630
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4629 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43962689

<400> 4630
tactgagggt ctatgcatgg taattagcag aaaacatgat agaactaaac a 51

<210> 4631
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4632 is other entry)

<221> misc_feature

<222> (0)...(0)
<223> Accession number cg43962735

<400> 4631
taactttctgc agtactttgt tcatataaaa cactagtaaa ataggcttct t 51

<210> 4632
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4631 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43962735

<400> 4632
taactttctgc agtactttgt tcataaaaaa cactagtaaa ataggcttct t 51

<210> 4633
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4634 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43962735

<400> 4633
taaaataggc ttcttaaaaa ttaaatagtg aaataccaac caaattatat a 51

<210> 4634
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4633 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43962735

<400> 4634
taaaataggc ttcttaaaaa ttaaacagtg aaataccaac caaattatat a 51

<210> 4635
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4636 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43962735

<400> 4635
aaccaaatta tatacattgt tacagtacaa gtgaatgagg caaaatatcc a 51

<210> 4636
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4635 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43962735

<400> 4636
aaccaaatta tatacattgt tacagcacia gtgaatgagg caaaatatcc a 51

<210> 4637
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4638 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43962735

<400> 4637
agtgcgtggc acggaggggg tgacaggaag gccacgttcc aatgtcacag t 51

<210> 4638
<211> 51
<212> DNA
<213> Homo sapiens

<220>

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<222> (26)...(0)
<223> 2 of 2 allelic variants (4637 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43962735

<400> 4638
agtgcgtggc acggaggggg tgacaagaag gccacgttcc aatgtcacag t 51

<210> 4639
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4640 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43962852

<400> 4639
taacaattta gcagccgtgg caactgcca ggacacatac aacaaaaaaaa t 51

<210> 4640
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4639 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43962852

<400> 4640
taacaattta gcagccgtgg caactaccaa ggacacatac aacaaaaaaaa t 51

<210> 4641
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4642 is other entry)

<221> misc_feature
<222> (0)...(0)

<223> Accession number cg43962852

<400> 4641

tggccccaaa tgacttcgag accaaacacc tgcaacttaa ggaagaatct g

51

<210> 4642

<211> 51

<212> DNA

<213> Homo sapiens

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<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (4641 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43962852

<400> 4642

tggccccaaa tgacttcgag accaagcacc tgcaacttaa ggaagaatct g

51

<210> 4643

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (4644 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43962884

<400> 4643

cagaaaatgc tttatttttc tctttgttcc ctcccatcc tatatttttc t

51

<210> 4644

<211> 50

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (4643 is other entry)

<221> misc_feature

<222> (25)...(26)

<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43962884

<400> 4644
cagaaaatgc tttatttttc tctttttccc tccccatcct atatttttct 50

<210> 4645
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4646 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43962884

<400> 4645
cctccccatc ctatatTTTT ctctaaaaa accctattat cagaaatatt a 51

<210> 4646
<211> 50
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4645 is other entry)

<221> misc_feature
<222> (25)...(26)
<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43962884

<400> 4646
cctccccatc ctatatTTTT ctctaaaaa ccctattatc agaaatatta 50

<210> 4647
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4648 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43963085

<400> 4647
gaccggtgcc acctctggaa tgggtgttcgc cgcgaaataa gctactggcg a 51

<210> 4648
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4647 is other entry)

<221> misc_feature
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<223> Accession number cg43963085

<400> 4648
gaccggtgcc acctctggaa tgggtgctcgc cgcgaaataa gctactggcg a 51

<210> 4649
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4650 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43963560

<400> 4649
ctgcttcagc ctcccaagta gctgagatta caggcaccca ccatcacgcc t 51

<210> 4650
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4649 is other entry)

<221> misc_feature
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<223> Accession number cg43963560

<400> 4650
ctgcttcagc ctcccaagta gctgaaatta caggcaccca ccatcacgcc t 51

<210> 4651
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4652 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43964079

<400> 4651
acccatccaa aattattttgt gataggtgaa aaatggccac aagctctttg t 51

<210> 4652
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4651 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43964079

<400> 4652
acccatccaa aattattttgt gatagatgaa aaatggccac aagctctttg t 51

<210> 4653
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4654 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43964913

<400> 4653
cgcctttctt tcccagcaga aagggatccg ttccggacag gacagaagtg a 51

<210> 4654
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4653 is other entry)

<221> misc_feature

<222> (0)...(0)
<223> Accession number cg43964913

<400> 4654
cgcttttctt tcccagcaga aaggggtccg ttccggacag gacagaagtg a 51

<210> 4655
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4656 is other entry)

<221> misc_feature
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<223> Accession number cg43964913

<400> 4655
agcagaaagg gatccgttcc ggacaggaca gaagtgagca gatggtttcc c 51

<210> 4656
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4655 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43964913

<400> 4656
agcagaaagg gatccgttcc ggacaagaca gaagtgagca gatggtttcc c 51

<210> 4657
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4658 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43964913

<400> 4657
gaaagggatc cggtccggac aggacagaag ttagcagatg gtttccccta c 51

<210> 4658
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4657 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43964913

<400> 4658
gaaagggatc cggtccggac aggaccgaag tgagcagatg gtttccccta c 51

<210> 4659
<211> 51
<212> DNA
<213> Homo sapiens

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<222> (26)...(0)
<223> 1 of 2 allelic variants (4660 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43964975

<400> 4659
cccaagttac tgcataccaa gcagctaata aaaaccaact gacttaaagt c 51

<210> 4660
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4659 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43964975

<400> 4660
cccaagttac tgcataccaa gcagccaata aaaaccaact gacttaaagt c 51

<210> 4661
<211> 51
<212> DNA
<213> Homo sapiens

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<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4662 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43965502

<400> 4661
acactcatta tactttttcc ccaaacaggt acaaaggagg ttcaagtgct c 51

<210> 4662
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (4661 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43965502

<400> 4662
acactcatta tactttttcc ccaaaaaggt acaaaggagg ttcaagtgct c 51

<210> 4663
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4664 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43966333

<400> 4663
ggttccgcac atcttttgaat cgctcccctt tgggtgctcac tacctggttg t 51

<210> 4664
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4663 is other entry)

<221> misc_feature
<222> (0)...(0)

<223> Accession number cg43966333

<400> 4664

ggttccgcac atctttgaat cgctctcctt tgggtgctcac tacctggttg t 51

<210> 4665

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 1 of 2 allelic variants (4666 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43966333

<400> 4665

atcgctcccc tttggtgctc actacctggt tgcgatgta tcggatcagc t 51

<210> 4666

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (4665 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43966333

<400> 4666

atcgctcccc tttggtgctc actacttgggt tgcgatgta tcggatcagc t 51

<210> 4667

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (4668 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43966536

<400> 4667

gcccttgccc tgggtggccc ggggccgcag cgcagagcag caggtgagca g 51

<210> 4668

<211> 50
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4667 is other entry)

<221> misc_feature
<222> (25)...(26)
<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43966536

<400> 4668
gcccctggcc tgggtggccc ggggcgcagc gcagagcagc aggtgagcag

50

<210> 4669
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4670 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43966551

<400> 4669
ggtggcttct cagcaccttg agcctctgat gcccgcctc tgacctcagg t

51

<210> 4670
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4669 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43966551

<400> 4670
ggtggcttct cagcaccttg agcctatgat gcccgcctc tgacctcagg t

51

<210> 4671
<211> 51
<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (4672 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43966619

<400> 4671

acacaggaag tgagttcaga tttatccaca aatgtcttct ggtctagttc t

51

<210> 4672

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (4671 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43966619

<400> 4672

acacaggaag tgagttcaga tttattcaca aatgtcttct ggtctagttc t

51

<210> 4673

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (4674 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43966787

<400> 4673

ctgttttagtg aaaacagtta agggcaggcc acttctacac tcccagcgct a

51

<210> 4674

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (4673 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43966787

<400> 4674
ctgttttagtg aaaacagtta agggccggcc acttctacac tcccagcgct a 51

<210> 4675
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4676 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43966806

<400> 4675
tatatacaca acattcaaga tattcttaat gtaagacatt tcagattgaa g 51

<210> 4676
<211> 50
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4675 is other entry)

<221> misc_feature
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<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43966806

<400> 4676
tatatacaca acattcaaga tattctaag taagacattt cagattgaag 50

<210> 4677
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4678 is other entry)

<221> misc_feature

<222> (0)...(0)
<223> Accession number cg43966806

<400> 4677
atatacacia cattcaagat attcttaatg taagacattt cagattgaag t 51

<210> 4678
<211> 50
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4677 is other entry)

<221> misc_feature
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<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43966806

<400> 4678
atatacacia cattcaagat attctaatgt aagacatttc agattgaagt 50

<210> 4679
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4680 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43966820

<400> 4679
caaactcgtc ataatgaac tgaagcgta gggctgactt gccaacgcct c 51

<210> 4680
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4679 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43966820

<400> 4680
caaactcgtc atacatgaac tgaagtgtca gggctgactt gccaacgcct c 51

<210> 4681
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4682 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43966864

<400> 4681
tctatggtaa atccttgcaa acatggaaac aatgcatttg gcccagtgc t 51

<210> 4682
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4681 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43966864

<400> 4682
tctatggtaa atccttgcaa acatgtaaac aatgcatttg gcccagtgc t 51

<210> 4683
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4684 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43967021

<400> 4683
tggaaaaaaa agagcagccc tctaaagagg gtgtcaaaca gtggaagaaa t 51

<210> 4684
<211> 51

<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4683 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43967021

<400> 4684
tggaaaaaaa agagcagccc tctaaggagg gtgtcaaaca gtggaagaaa t 51

<210> 4685
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4686 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43967079

<400> 4685
atgacccctt actacacaca cacacacgca aagttagggtc cagcataggc c 51

<210> 4686
<211> 50
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4685 is other entry)

<221> misc_feature
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<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
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<223> Accession number cg43967079

<400> 4686
atgacccctt actacacaca cacaccgcaa agttagggtcc agcataggcc 50

<210> 4687
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4688 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43967102

<400> 4687
tccatcaccc tgctggggct cgccgtcaac gtggtcacca cgctcgtgct c 51

<210> 4688
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4687 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43967102

<400> 4688
tccatcaccc tgctggggct cgccgccaac gtggtcacca cgctcgtgct c 51

<210> 4689
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4690 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43967102

<400> 4689
ccacagtatt tatggcagtg ggagcttcaa ttgccgctcg cttaggaact t 51

<210> 4690
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4689 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43967102

<400> 4690
ccacagtatt tatggcagtg ggagcctcaa ttgccgctcg cttaggaact t 51

<210> 4691
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4692 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43967119

<400> 4691
gcagggcaaa gaccagacta tctccacctg ttgttttgca ttgtttctgg g 51

<210> 4692
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4691 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43967119

<400> 4692
gcagggcaaa gaccagacta tctccgctg ttgttttgca ttgtttctgg g 51

<210> 4693
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4694 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43967630

<400> 4693
agctgctgtc cccagagagg agacaacagc ttctggaggc tctggggact c 51

<210> 4694
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4693 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43967630

<400> 4694
agctgctgtc cccagagagg agacagcagc ttctggaggc tctggggact c 51

<210> 4695
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4696 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43967706

<400> 4695
cactgggcaa ggtaggtagc tagctgcctg acccctagtc tggggttgga a 51

<210> 4696
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4695 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43967706

<400> 4696
cactgggcaa ggtaggtagc tagcttctctg acccctagtc tggggttgga a 51

<210> 4697
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4698 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43967833

<400> 4697
aaacatatca aaatgttaca aaaatgtatg gctcccttgc tgaggccctg t 51

<210> 4698
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4697 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43967833

<400> 4698
aaacatatca aaatgttaca aaaatatatg gctcccttgc tgaggccctg t 51

<210> 4699
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4700 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43967844

<400> 4699
acaactaata atcttctttc aagagttttt ttttcaatc ttgaggttaa c 51

<210> 4700
<211> 50
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4699 is other entry)

<221> misc_feature

<222> (25)...(26)
<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43967844

<400> 4700
acaactaata atcttcttttc aagagttttt ttttcaatct tggagttaac 50

<210> 4701
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4702 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43967844

<400> 4701
tcatgcttct gggttgaaag tgacgagtaa atatgtcaga ctgtttaaag g 51

<210> 4702
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4701 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43967844

<400> 4702
tcatgcttct gggttgaaag tgacgggtaa atatgtcaga ctgtttaaag g 51

<210> 4703
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4704 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43968063

<400> 4703
gcggctcgat acgcagtgag aagtcaccgc gctcaaagggc atccgcgccc a 51

<210> 4704
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4703 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43968063

<400> 4704
gcggctcgat acgcagtgag aagtccccgc gctcaaagggc atccgcgccc a 51

<210> 4705
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4706 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43968079

<400> 4705
ctgtgagcgt gtctgatgcc ccgaacaggt gccagggtccc ccaaaagcag c 51

<210> 4706
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4705 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43968079

<400> 4706
ctgtgagcgt gtctgatgcc ccgaaaaggt gccagggtccc ccaaaagcag c 51

<210> 4707
<211> 51

<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4708 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43968177

<400> 4707
aaatgctgtc cagttttatt ttttttatgt ttttatcctt ggatgtacaa a 51

<210> 4708
<211> 50
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4707 is other entry)

<221> misc_feature
<222> (25)...(26)
<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43968177

<400> 4708
aaatgctgtc cagttttatt tttttatggt gttatccttg gatgtacaaa 50

<210> 4709
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4710 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43968179

<400> 4709
gaaattcttc agagatcttc aaagcacaaa aaatacgttc ttttttcaaa g 51

<210> 4710
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4709 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43968179

<400> 4710
gaaattcttc agagatcttc aaagcgcaaa aaatacgttc ttttttcaaa g 51

<210> 4711
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4712 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43968211

<400> 4711
tggaaaatgt agttggagat aaagtttttg gaagctttgc tgaaaacctt t 51

<210> 4712
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4711 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43968211

<400> 4712
tggaaaatgt agttggagat aaagtctttg gaagctttgc tgaaaacctt t 51

<210> 4713
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4714 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43968211

<400> 4713
ttggaagctt tgctgaaaac ctttcatttc ttctggaagc tttaaaaaaa g 51

<210> 4714
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4713 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43968211

<400> 4714
ttggaagctt tgctgaaaac ctttcgtttc ttctggaagc tttaaaaaaa g 51

<210> 4715
<211> 51
<212> DNA
<213> Homo sapiens

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<222> (26)...(0)
<223> 1 of 2 allelic variants (4716 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43968211

<400> 4715
tcttctggaa gctttaaaaa aaggtgaccg aactagcagt tgcccagtga t 51

<210> 4716
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (4715 is other entry)

<221> misc_feature
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<223> Accession number cg43968211

<400> 4716
tcttctggaa gctttaaaaa aaggtgaccg aactagcagt tgcccagtga t 51

<210> 4717
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4718 is other entry)

<221> misc_feature
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<223> Accession number cg43968211

<400> 4717
tggaagcttt aaaaaaaggt gaccgaacta gcagttgccc agtgatcttc a 51

<210> 4718
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (4717 is other entry)

<221> misc_feature
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<223> Accession number cg43968211

<400> 4718
tggaagcttt aaaaaaaggt gaccggacta gcagttgccc agtgatcttc a 51

<210> 4719
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (4720 is other entry)

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<223> Accession number cg43968211

<400> 4719
tagcagttgc ccagtgatct tcatattaga tgaatttgat ctttttgc a 51

<210> 4720
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (4719 is other entry)

<221> misc_feature
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<223> Accession number cg43968211

<400> 4720
tagcagttgc ccagtgatct tcatactaga tgaatttgat ctttttgctc a 51

<210> 4721
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (4722 is other entry)

<221> misc_feature
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<223> Accession number cg43968211

<400> 4721
ttgctcatca taaaaaccaa acacttctct ataatctttt tgacatttct c 51

<210> 4722
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (4721 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43968211

<400> 4722
ttgctcatca taaaaaccaa acactcctct ataatctttt tgacatttct c 51

<210> 4723
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (4724 is other entry)

<221> misc_feature

<222> (0)...(0)
<223> Accession number cg43968211

<400> 4723
tctataatct ttttgacatt tctcagtctg cacagacccc aatagcagtt a 51

<210> 4724
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4723 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43968211

<400> 4724
tctataatct ttttgacatt tctcaatctg cacagacccc aatagcagtt a 51

<210> 4725
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4726 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43968211

<400> 4725
cacagacccc aatagcagtt attggtctta catgtagatt ggatattttg g 51

<210> 4726
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4725 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43968211

<400> 4726
cacagacccc aatagcagtt attggcctta catgtagatt ggatattttg g 51

<210> 4727
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4728 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43968242

<400> 4727
gcgtactggc gacccggagt gatgagcccg cccgagacga tgccgccgtg g 51

<210> 4728
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4727 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43968242

<400> 4728
gcgtactggc gacccggagt gatgaccccg cccgagacga tgccgccgtg g 51

<210> 4729
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4730 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43968298

<400> 4729
ttgacaattt tggtttgggg cttttgcttc ggtccaagca gataaaacgc a 51

<210> 4730
<211> 51
<212> DNA
<213> Homo sapiens

<220>

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<223> 2 of 2 allelic variants (4729 is other entry)

<221> misc_feature
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<223> Accession number cg43968298

<400> 4730
ttgacaattt tggtttgggg cttttacttc ggtccaagca gataaaaacgc a 51

<210> 4731
<211> 51
<212> DNA
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<220>
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<223> 1 of 2 allelic variants (4732 is other entry)

<221> misc_feature
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<223> Accession number cg43968298

<400> 4731
attttggttt ggggcttttg cttcgggtcca agcagataaa acgcatgggc t 51

<210> 4732
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 2 of 2 allelic variants (4731 is other entry)

<221> misc_feature
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<400> 4732
attttggttt ggggcttttg cttcgatcca agcagataaa acgcatgggc t 51

<210> 4733
<211> 51
<212> DNA
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<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4734 is other entry)

<221> misc_feature
<222> (0)...(0)

<223> Accession number cg43968308

<400> 4733

tctcagcctc ggggctgcaa tccagggctg tgctgagcag caagaggaga g

51

<210> 4734

<211> 51

<212> DNA

<213> Homo sapiens

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<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (4733 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43968308

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tctcagcctc ggggctgcaa tccagagctg tgctgagcag caagaggaga g

51

<210> 4735

<211> 51

<212> DNA

<213> Homo sapiens

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<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (4736 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43968538

<400> 4735

tctgttggtta atcggttaca ttgtcacctc taataccagt catcaaatcc a

51

<210> 4736

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (4735 is other entry)

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<222> (0)...(0)

<223> Accession number cg43968538

<400> 4736

tctgttggtta atcggttaca ttgtctctc taataccagt catcaaatcc a

51

<210> 4737

<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4738 is other entry)

<221> misc_feature
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<223> Accession number cg43968538

<400> 4737
tcaggtcata ggattccttt ttttttaaag ataagtaa at gcattccagaa a 51

<210> 4738
<211> 50
<212> DNA
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4737 is other entry)

<221> misc_feature
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<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43968538

<400> 4738
tcaggtcata ggattccttt tttttaaaga taagtaa atg catccagaaa 50

<210> 4739
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4740 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43968629

<400> 4739
tggctctggc tggcttccat ggggggctca tcaactggaag ggctgggtgac.c 51

<210> 4740
<211> 51
<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (4739 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43968629

<400> 4740

tggtctggc tggcttccat gggggactca tcaactggaag ggctggtgac c

51

<210> 4741

<211> 51

<212> DNA

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<223> 1 of 2 allelic variants (4742 is other entry)

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<222> (0)...(0)

<223> Accession number cg43968904

<400> 4741

ccatagcaat tagcatat t tctaacaagc catgttggtt aaatttatat t

51

<210> 4742

<211> 51

<212> DNA

<213> Homo sapiens

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<222> (26)...(0)

<223> 2 of 2 allelic variants (4741 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43968904

<400> 4742

ccatagcaat tagcatat t tctaataagc catgttggtt aaatttatat t

51

<210> 4743

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<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (4744 is other entry)

<221> misc_feature
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<223> Accession number cg43968904

<400> 4743
tgaagagtat gtgatatatt gtcaaggggtg taaagaaaac tctccctggt t 51

<210> 4744
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4743 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43968904

<400> 4744
tgaagagtat gtgatatatt gtcaatgggtg taaagaaaac tctccctggt t 51

<210> 4745
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4746 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43969016

<400> 4745
acgatcatat tttggcgtga agaagagggc aagagaagca gaaggagaat g 51

<210> 4746
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4745 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43969016

<400> 4746

acgatcatat tttggcgtga agaaggggggc aagagaagca gaaggagaat g

51

<210> 4747

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (4748 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43969342

<400> 4747

cggtcactg catcctccgc ctcccaggtt caagctattc tctgcctca g

51

<210> 4748

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (4747 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43969342

<400> 4748

cggtcactg catcctccgc ctcccgggtt caagctattc tctgcctca g

51

<210> 4749

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 1 of 2 allelic variants (4750 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43969342

<400> 4749

acggggtttc accatgttgg ccaggctagt ttcgaactcc tgacctcagt g

51

<210> 4750

<211> 51

<212> DNA

<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4749 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43969342

<400> 4750
acgggggtttc accatgttgg ccaggatagt ttcgaactcc tgacctcagt g 51

<210> 4751
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4752 is other entry)

<221> misc_feature
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<223> Accession number cg43969665

<400> 4751
agacagaatg ggggaaaatg gggagaaaaa aaaacaaacc ttccttttcc c 51

<210> 4752
<211> 50
<212> DNA
<213> Homo sapiens

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<222> (26)...(0)
<223> 2 of 2 allelic variants (4751 is other entry)

<221> misc_feature
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<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43969665

<400> 4752
agacagaatg ggggaaaatg gggagaaaaa aaacaaacct tcccttttccc 50

<210> 4753
<211> 51
<212> DNA
<213> Homo sapiens

<220>

<221> misc_feature
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<223> 1 of 2 allelic variants (4754 is other entry)

<221> misc_feature
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<223> Accession number cg43969665

<400> 4753
ctgggagacc atgtggtata aaaaaagtca ttaaagttgc tgcagaaaa a 51

<210> 4754
<211> 50
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4753 is other entry)

<221> misc_feature
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<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43969665

<400> 4754
ctgggagacc atgtggtata aaaaagtcac taaagttgct tgcagaaaaa 50

<210> 4755
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4756 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43970196

<400> 4755
gacatcccaa actttacaaa aacttacaat gctgcttata taaaactttt c 51

<210> 4756
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (4755 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43970196

<400> 4756

gacatcccaa actttacaaa aacttgcaat gctgcttatc taaaactttt c

51

<210> 4757

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (4758 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43970375

<400> 4757

acaggtaagc ggggaaacgt acctatgact ctggcaaaat attctcggca t

51

<210> 4758

<211> 51

<212> DNA

<213> Homo sapiens

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<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (4757 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43970375

<400> 4758

acaggtaagc ggggaaacgt acctaggact ctggcaaaat attctcggca t

51

<210> 4759

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (4760 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43970424

<400> 4759
aaggtgcctg cctgcctggg gcctaacgag ccaaacagtg ctaatttcat c 51

<210> 4760
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4759 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43970424

<400> 4760
aaggtgcctg cctgcctggg gcctaccgag ccaaacagtg ctaatttcat c 51

<210> 4761
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4762 is other entry)

<221> misc_feature
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<223> Accession number cg43970424

<400> 4761
gtgcctgcct gcctggggcc taacgagcca aacagtgcta atttcatcca t 51

<210> 4762
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4761 is other entry)

<221> misc_feature
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<223> Accession number cg43970424

<400> 4762
gtgcctgcct gcctggggcc taacgcgcca aacagtgcta atttcatcca t 51

<210> 4763
<211> 51
<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (4764 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43970474

<400> 4763

tctatttccc tcaaaagttt ttttttcctg ctataagata aagaaaaggc t

51

<210> 4764

<211> 50

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (4763 is other entry)

<221> misc_feature

<222> (25)...(26)

<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43970474

<400> 4764

tctatttccc tcaaaagttt tttttcctgc tataagataa agaaaaggct

50

<210> 4765

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (4766 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43970521

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gcccacagcc tcaggaacag cccagactgg ggaggcgtgg tcacactaca c

51

<210> 4766

<211> 51

<212> DNA

<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (4765 is other entry)

<221> misc_feature
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<223> Accession number cg43970521

<400> 4766
gccacagcc tcaggaacag cccagcctgg ggaggcgtgg tcacactaca c 51

<210> 4767
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4768 is other entry)

<221> misc_feature
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<223> Accession number cg43970521

<400> 4767
ttaactgtgg gcgatgagaa aggtggtgga ctcttcttac tggcagggcc a 51

<210> 4768
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (4767 is other entry)

<221> misc_feature
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<223> Accession number cg43970521

<400> 4768
ttaactgtgg gcgatgagaa aggtgctgga ctcttcttac tggcagggcc a 51

<210> 4769
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4770 is other entry)

<221> misc_feature

<222> (0)...(0)
<223> Accession number cg43970521

<400> 4769
cttcttactg gcagggccac ctgcgtctgt ggagaccctg gggcccaggg t 51

<210> 4770
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4769 is other entry)

<221> misc_feature
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<223> Accession number cg43970521

<400> 4770
cttcttactg gcagggccac ctgcggctgt ggagaccctg gggcccaggg t 51

<210> 4771
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (4772 is other entry)

<221> misc_feature
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<223> Accession number cg43970716

<400> 4771
cgtggaggct tcgttttcta tttatttaca ttattggctt tcttttgtga g 51

<210> 4772
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 2 of 2 allelic variants (4771 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43970716

<400> 4772
cgtggaggct tcgttttcta tttatctaca ttattggctt tcttttgtga g 51

<210> 4773
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 1 of 2 allelic variants (4774 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43970716

<400> 4773
tctgccgact ccaaggtagg gatggggctg tccccaacag acaccagcg a 51

<210> 4774
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4773 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43970716

<400> 4774
tctgccgact ccaaggtagg gatggagctg tccccaacag acaccagcg a 51

<210> 4775
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 1 of 2 allelic variants (4776 is other entry)

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<223> Accession number cg43970716

<400> 4775
ccgactccaa ggtagggatg gggctgtccc caacagacac cagcgacat g 51

<210> 4776
<211> 50
<212> DNA
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<223> 2 of 2 allelic variants (4775 is other entry)

<221> misc_feature
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<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
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<400> 4776
ccgactccaa ggtagggatg gggcttcccc aacagacacc agcgacatg 50

<210> 4777
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 1 of 2 allelic variants (4778 is other entry)

<221> misc_feature
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<223> Accession number cg43970716

<400> 4777
agggatgggg ctgtccccaa cagacaccag cgcacatgcc ctatttggtat 51

<210> 4778
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (4777 is other entry)

<221> misc_feature
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<223> Accession number cg43970716

<400> 4778
agggatgggg ctgtccccaa cagaccccag cgcacatgcc ctatttggtat 51

<210> 4779
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 1 of 2 allelic variants (4780 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43970764

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caacgtccgt gtcgatgccg aaatcctttt ttaaactctt ttttgaggga a

51

<210> 4780

<211> 51

<212> DNA

<213> Homo sapiens

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<222> (26)...(0)

<223> 2 of 2 allelic variants (4779 is other entry)

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<223> Accession number cg43970764

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caacgtccgt gtcgatgccg aaatcctttt ttaaactctt ttttgaggga a

51

<210> 4781

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 1 of 2 allelic variants (4782 is other entry)

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<223> Accession number cg43970764

<400> 4781

ctctatatttg tttcccatct tcctcctgt tctctcccat cctccaaaga t

51

<210> 4782

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 2 of 2 allelic variants (4781 is other entry)

<221> misc_feature

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<223> Accession number cg43970764

<400> 4782
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<210> 4783
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (4784 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43970806

<400> 4783
aaactaaaga tttgtgtgcc gcacggtgat tctgccctc tggccttct t 51

<210> 4784
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 2 of 2 allelic variants (4783 is other entry)

<221> misc_feature
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<223> Accession number cg43970806

<400> 4784
aaactaaaga tttgtgtgcc gcacgttgat tctgccctc tggccttct t 51

<210> 4785
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4786 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43970995

<400> 4785
taccaggaaa ctgttacaga cgccattttt ttttttttg agacggagtc t 51

<210> 4786
<211> 50
<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (4785 is other entry)

<221> misc_feature

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<223> Nucleotide deleted between bases 25 and 26

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<222> (0)...(0)

<223> Accession number cg43970995

<400> 4786

taccaggaaa ctgttacaga cgccattttt ttttttttga gacggagtct

50

<210> 4787

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 1 of 2 allelic variants (4788 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43970995

<400> 4787

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51

<210> 4788

<211> 50

<212> DNA

<213> Homo sapiens

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<223> 2 of 2 allelic variants (4787 is other entry)

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<222> (25)...(26)

<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature

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<223> Accession number cg43970995

<400> 4788

gttacagacg ccattttttt ttttttgaga cggagtcttg ctctgttgcc

50

<210> 4789

<211> 51
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<223> 1 of 2 allelic variants (4790 is other entry)

<221> misc_feature
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<223> Accession number cg43970995

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ttacagacgc catttttttt ttttttgaga cggagtcttg ctctgttgcc c 51

<210> 4790
<211> 50
<212> DNA
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<223> 2 of 2 allelic variants (4789 is other entry)

<221> misc_feature
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<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
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<223> Accession number cg43970995

<400> 4790
ttacagacgc catttttttt tttttgagac ggagtcttgc tctgttgccc 50

<210> 4791
<211> 51
<212> DNA
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<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4792 is other entry)

<221> misc_feature
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<223> Accession number cg43971398

<400> 4791
taatagtaga tttttaatca gctttgtgat ttctttaata gttattgggt t 51

<210> 4792
<211> 51
<212> DNA

<213> Homo sapiens

<220>

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<222> (26)...(0)

<223> 2 of 2 allelic variants (4791 is other entry)

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<223> Accession number cg43971398

<400> 4792

taatagtaga tttttaatca gctttctgat ttctttaata gttattggtt t

51

<210> 4793

<211> 51

<212> DNA

<213> Homo sapiens

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<222> (26)...(0)

<223> 1 of 2 allelic variants (4794 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43971504

<400> 4793

atggccggga catctcagag cacacgcatg accaggtggt gatgttcac a

51

<210> 4794

<211> 51

<212> DNA

<213> Homo sapiens

<220>

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<222> (26)...(0)

<223> 2 of 2 allelic variants (4793 is other entry)

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<222> (0)...(0)

<223> Accession number cg43971504

<400> 4794

atggccggga catctcagag cacactcatg accaggtggt gatgttcac a

51

<210> 4795

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (4796 is other entry)

<221> misc_feature
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<223> Accession number cg43971504

<400> 4795
tcaaagcaag ccgggagtc cactcaagag aactggccct ggtgatcagg a 51

<210> 4796
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (4795 is other entry)

<221> misc_feature
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<223> Accession number cg43971504

<400> 4796
tcaaagcaag ccgggagtc cactcgagag aactggccct ggtgatcagg a 51

<210> 4797
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (4798 is other entry)

<221> misc_feature
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<223> Accession number cg43971504

<400> 4797
gagaactggc cctggtgatc aggaggagag ctgtgcgctc atttgctgac t 51

<210> 4798
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (4797 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43971504

<400> 4798

gagaactggc cctggtgatc aggagaagag ctgtgcgctc atttgctgac t 51

<210> 4799

<211> 51

<212> DNA

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<210> 4800

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<212> DNA

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<210> 4801

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 1 of 2 allelic variants (4802 is other entry)

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<223> Accession number cg43971504

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<210> 4802

<211> 51

<212> DNA

<213> Homo sapiens

11.17

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<221> misc_feature
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<223> Accession number cg43971504

<400> 4802
tgatcaggag gagagctgtg cgctcgtttg ctgacttcaa gtctgaagat g 51

<210> 4803
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<212> DNA
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<223> 1 of 2 allelic variants (4804 is other entry)

<221> misc_feature
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<223> Accession number cg43971504

<400> 4803
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<210> 4804
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 2 of 2 allelic variants (4803 is other entry)

<221> misc_feature
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<400> 4804
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<210> 4805
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<212> DNA
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<223> 1 of 2 allelic variants (4806 is other entry)

<221> misc_feature
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<400> 4805
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<210> 4806
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
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<223> 2 of 2 allelic variants (4805 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43971504

<400> 4806
ttgctgactt caagtctgaa gatgagctga accagctttt cccggaagcc a 51

<210> 4807
<211> 51
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<223> 1 of 2 allelic variants (4808 is other entry)

<221> misc_feature
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<400> 4807
gacttcaagt ctgaagatga actgaaccag cttttcccg aagccatttt c 51

<210> 4808
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (4807 is other entry)

<221> misc_feature
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<400> 4808
gacttcaagt ctgaagatga actgagccag cttttcccg aagccatttt c 51

<210> 4809
<211> 51
<212> DNA
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<220>
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<223> 1 of 2 allelic variants (4810 is other entry)

<221> misc_feature
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<400> 4809
aagatgaact gaaccagctt ttccccgaag ccattttccc catgtgtccg g 51

<210> 4810
<211> 51
<212> DNA
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<223> 2 of 2 allelic variants (4809 is other entry)

<221> misc_feature
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<400> 4810
aagatgaact gaaccagctt ttcccagaag ccattttccc catgtgtccg g 51

<210> 4811
<211> 51
<212> DNA
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<220>
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<223> 1 of 2 allelic variants (4812 is other entry)

<221> misc_feature
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<223> Accession number cg43971504

<400> 4811
tgaaccagct tttccccgaa gccattttcc ccatgtgtcc ggagggtggg g 51

<210> 4812
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (4811 is other entry)

<221> misc_feature
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<223> Accession number cg43971504

<400> 4812
tgaaccagct tttccccgaa gccatgttcc ccatgtgtcc ggagggtggg g 51

<210> 4813
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4814 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43971504

<400> 4813
cccatttacc cactggatat tgtccgaaaa atgcgagacc agcgcgccat g 51

<210> 4814
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4813 is other entry)

<221> misc_feature
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<400> 4814
cccatttacc cactggatat tgtccaaaaa atgcgagacc agcgcgccat g 51

<210> 4815
<211> 51
<212> DNA
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<220>
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<223> 1 of 2 allelic variants (4816 is other entry)

<221> misc_feature

<222> (0)...(0)
<223> Accession number cg43971764

<400> 4815
aattttttgt atttttagta gagacgggtt tcaccgtgtt agccaggatg g 51

<210> 4816
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 2 of 2 allelic variants (4815 is other entry)

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<400> 4816
aattttttgt atttttagta gagacagggtt tcaccgtgtt agccaggatg g 51

<210> 4817
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<212> DNA
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<223> 1 of 2 allelic variants (4818 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43971764

<400> 4817
caggatggtc tcgatctcct gaccttgtga tccgcccgcc tcagcctccc a 51

<210> 4818
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4817 is other entry)

<221> misc_feature
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<223> Accession number cg43971764

<400> 4818
caggatggtc tcgatctcct gacctcgtga tccgcccgcc tcagcctccc a 51

<210> 4819
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<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4820 is other entry)

<221> misc_feature
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<223> Accession number cg43972205

<400> 4819
ggtcaccaag atgcagcagg aaatcacttt tcagcaagta atgtctcaga t 51

<210> 4820
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4819 is other entry)

<221> misc_feature
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<400> 4820
ggtcaccaag atgcagcagg aaatcgcttt tcagcaagta atgtctcaga t 51

<210> 4821
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4822 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43972259

<400> 4821
ttttagtgtc aaaatatagc gttgagggga gctggacgct agggctcttca c 51

<210> 4822
<211> 51
<212> DNA
<213> Homo sapiens

<220>

<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4821 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43972259

<400> 4822
tttttagtgc aaaatatagc gttgaaggga gctggacgct agggctcttca c 51

<210> 4823
<211> 51
<212> DNA
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<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4824 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43972293

<400> 4823
tgacattaaa gaagaaacag acaccttgga gaatttatga ctcctttctc t 51

<210> 4824
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4823 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43972293

<400> 4824
tgacattaaa gaagaaacag acaccctgga gaatttatga ctcctttctc t 51

<210> 4825
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4826 is other entry)

<221> misc_feature
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<223> Accession number cg43972406

<400> 4825

gtgattccat ctgaaggcag tgagaagtgg tgttaccatc tgaagtgggt c

51

<210> 4826

<211> 51

<212> DNA

<213> Homo sapiens

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<222> (26)...(0)

<223> 2 of 2 allelic variants (4825 is other entry)

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<223> Accession number cg43972406

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gtgattccat ctgaaggcag tgagaggtgg tgttaccatc tgaagtgggt c

51

<210> 4827

<211> 51

<212> DNA

<213> Homo sapiens

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<222> (26)...(0)

<223> 1 of 2 allelic variants (4828 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43972482

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51

<210> 4828

<211> 51

<212> DNA

<213> Homo sapiens

<220>

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<222> (26)...(0)

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<223> Accession number cg43972482

<400> 4828

ctctggctcc ggagtagctg ggattacagg caccgccac cagcctggc t

51

<210> 4829

<211> 51
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4830 is other entry)

<221> misc_feature
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<223> Accession number cg43972482

<400> 4829
gattgcaggc acccgccacc acgcctggct aatttttgta ttttagtag a 51

<210> 4830
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 2 of 2 allelic variants (4829 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43972482

<400> 4830
gattgcaggc acccgccacc acgcccggct aatttttgta ttttagtag a 51

<210> 4831
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 1 of 2 allelic variants (4832 is other entry)

<221> misc_feature
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<223> Accession number cg43972482

<400> 4831
cccgccacca cgcctggcta atttttgat ttttagtaga gacggggttt c 51

<210> 4832
<211> 50
<212> DNA
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<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43972482

<400> 4832
ccgccacca cgctggcta attttgtatt tttagtagag acggggtttc 50

<210> 4833
<211> 51
<212> DNA
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<223> 1 of 2 allelic variants (4834 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43972723

<400> 4833
gtggtaagtc tgggagtgga ggaaacaact ggtgtctgaa tatgactaaa a 51

<210> 4834
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 2 of 2 allelic variants (4833 is other entry)

<221> misc_feature
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<223> Accession number cg43972723

<400> 4834
gtggtaagtc tgggagtgga ggaaaaaact ggtgtctgaa tatgactaaa a 51

<210> 4835
<211> 51
<212> DNA
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<223> 1 of 2 allelic variants (4836 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43972879

<400> 4835
actgctgcag atccttagca ccattcagga actgtggagg gggcagcggg g 51

<210> 4836
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (4835 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43972879

<400> 4836
actgctgcag atccttagca ccatttagga actgtggagg gggcagcggg g 51

<210> 4837
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (4838 is other entry)

<221> misc_feature
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<223> Accession number cg43973078

<400> 4837
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<210> 4838
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4837 is other entry)

<221> misc_feature
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<223> Accession number cg43973078

<400> 4838

accttagaaa ggactcatta ccttgctgat atgggttggt tctatgtccc c

51

<210> 4839

<211> 51

<212> DNA

<213> Homo sapiens

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<222> (26)...(0)

<223> 1 of 2 allelic variants (4840 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43973078

<400> 4839

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51

<210> 4840

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 2 of 2 allelic variants (4839 is other entry)

<221> misc_feature

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<223> Accession number cg43973078

<400> 4840

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51

<210> 4841

<211> 51

<212> DNA

<213> Homo sapiens

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<222> (26)...(0)

<223> 1 of 2 allelic variants (4842 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43973078

<400> 4841

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51

<210> 4842

<211> 50

<212> DNA

<213> Homo sapiens

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<223> 2 of 2 allelic variants (4841 is other entry)

<221> misc_feature
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<221> misc_feature
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<223> Accession number cg43973078

<400> 4842
tcattacctt ggtgatatgg ttggtctat gtccccaccc aaatctcata 50

<210> 4843
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4844 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43973078

<400> 4843
cattaccttg gtgatatggt ttggtctat gtccccaccc aaatctcata t 51

<210> 4844
<211> 50
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4843 is other entry)

<221> misc_feature
<222> (25)...(26)
<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
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<223> Accession number cg43973078

<400> 4844
cattaccttg gtgatatggt ttggtctatg tccccaccca aatctcatat 50

<210> 4845
<211> 51

<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (4846 is other entry)

<221> misc_feature
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<223> Accession number cg43973078

<400> 4845
tgtccccacc caaatctcat atcgaattgt aatccccata atcccccatg t 51

<210> 4846
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4845 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43973078

<400> 4846
tgtccccacc caaatctcat atcgacttgt aatccccata atcccccatg t 51

<210> 4847
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4848 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43973078

<400> 4847
aatcccccat gttgaggag ggaccatatg ggaggtgact ggatcatggg a 51

<210> 4848
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 2 of 2 allelic variants (4847 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43973078

<400> 4848

aatcccccat gttgagggag ggaccttatg ggaggtgact ggatcatggg a

51

<210> 4849

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (4850 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43973114

<400> 4849

taattaaaca aacttaaaaa aaaaaaatag cattgggggac cctattttgt g

51

<210> 4850

<211> 50

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

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<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43973114

<400> 4850

taattaaaca aacttaaaaa aaaaaaatagc attgggggac ctattttgtg

50

<210> 4851

<211> 51

<212> DNA

<213> Homo sapiens

<220>

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<222> (26)...(0)

<223> 1 of 2 allelic variants (4852 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43973114

<400> 4851
aattaaacaa acttaaaaaa aaaaaatagc attgggggcc ctattttgtg a 51

<210> 4852
<211> 50
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (4851 is other entry)

<221> misc_feature
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<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43973114

<400> 4852
aattaaacaa acttaaaaaa aaaaaatagca ttgggggccc tatttttgtga 50

<210> 4853
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4854 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43973149

<400> 4853
atctacttac agtcttagta tgaaagtgtt cgggggtcct tgtaggttt g 51

<210> 4854
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4853 is other entry)

<221> misc_feature
<222> (0)...(0)

<223> Accession number cg43973149

<400> 4854

atctacttac agtcttagta tgaaaatggt cgggggctct tgtaggttt g

51

<210> 4855

<211> 51

<212> DNA

<213> Homo sapiens

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<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (4856 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43973149

<400> 4855

ctaatagtca tgcaaagtct taagcaaaaa agaagttaca ttaagcagaa c

51

<210> 4856

<211> 50

<212> DNA

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<223> 2 of 2 allelic variants (4855 is other entry)

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<221> misc_feature

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<223> Accession number cg43973149

<400> 4856

ctaatagtca tgcaaagtct taagcaaaaa gaagttacat taagcagaac

50

<210> 4857

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 1 of 2 allelic variants (4858 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43973149

<400> 4857
aatcaatgct taaaaaaciaa aaaaaacctg ggcagttcct aactacttaa a 51

<210> 4858
<211> 50
<212> DNA
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<223> 2 of 2 allelic variants (4857 is other entry)

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<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43973149

<400> 4858
aatcaatgct taaaaaaciaa aaaaacctgg gcagttccta actacttaaa 50

<210> 4859
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4860 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43973267

<400> 4859
gaagtgggtgg gaacaatgag ggacagcctg gatcatgtgg accagccaat g 51

<210> 4860
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (4859 is other entry)

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<223> Accession number cg43973267

<400> 4860
gaagtgggtgg gaacaatgag ggacaccctg gatcatgtgg accagccaat g 51

<210> 4861
<211> 51
<212> DNA
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<223> 1 of 2 allelic variants (4862 is other entry)

<221> misc_feature
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<223> Accession number cg43973267

<400> 4861
aagtgggtggg aacaatgagg gacagcctgg atcatgtgga ccagccaatg c 51

<210> 4862
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 2 of 2 allelic variants (4861 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43973267

<400> 4862
aagtgggtggg aacaatgagg gacaggctgg atcatgtgga ccagccaatg c 51

<210> 4863
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (4864 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43973459

<400> 4863
attttttatt aaaaattgat cagaagctag ttgaaattct caatgtaaat a 51

<210> 4864
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
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<223> 2 of 2 allelic variants (4863 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43973459

<400> 4864
atTTTTtatt aaaaattgat cagaaactag ttgaaattct caatgtaaat a 51

<210> 4865
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4866 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43973531

<400> 4865
aactggcagc cactattaat tgatcatcaa ctcaagctca agttgctgaa a 51

<210> 4866
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4865 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43973531

<400> 4866
aactggcagc cactattaat tgatcgtaa ctcaagctca agttgctgaa a 51

<210> 4867
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4868 is other entry)

<221> misc_feature

<222> (0)...(0)
<223> Accession number cg43973531

<400> 4867
acgcagccag acctagcttg ctaatgcaag gagaaaaggg gcaagttctt t 51

<210> 4868
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4867 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43973531

<400> 4868
acgcagccag acctagcttg ctaatacaag gagaaaaggg gcaagttctt t 51

<210> 4869
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4870 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43973700

<400> 4869
ctcaataatc tggtccttct caatgtccaa tctgtcatc tccaggtcca c 51

<210> 4870
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (4869 is other entry)

<221> misc_feature
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<223> Accession number cg43973700

<400> 4870
ctcaataatc tggtccttct caatgccc aa tctgtcatc tccaggtcca c 51

<210> 4871
<211> 50
<212> DNA
<213> Homo sapiens

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<223> 1 of 2 allelic variants (4872 is other entry)

<221> misc_feature
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<221> misc_feature
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<223> Accession number cg43973789

<400> 4871
ttccaattgt cctttttttt tttttaagag aattctggta tgagaaccat 50

<210> 4872
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4871 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43973789

<400> 4872
ttccaattgt cctttttttt tttttaaga gaattctggat atgagaacca t 51

<210> 4873
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4874 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43973789

<400> 4873
aaaaatttta ttctctgtcc catttatttg gtatctggaa cgaagagaaa a 51

<210> 4874
<211> 51

<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4873 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43973789

<400> 4874
aaaaatttta ttctctgtcc catttttttg gtatctggaa cgaagagaaa a 51

<210> 4875
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4876 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43974402

<400> 4875
ctgggattac aggcattgcac caccacaccc ggctaacttt tgtattttta g 51

<210> 4876
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4875 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43974402

<400> 4876
ctgggattac aggcattgcac caccataccc ggctaacttt tgtattttta g 51

<210> 4877
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)

<223> 1 of 2 allelic variants (4878 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43975023

<400> 4877

ttgataaaca acctcaaaga caccctcgagc ctttgaacgg agccctctgc a

51

<210> 4878

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (4877 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43975023

<400> 4878

ttgataaaca acctcaaaga caccctgagc ctttgaacgg agccctctgc a

51

<210> 4879

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (4880 is other entry)

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<222> (0)...(0)

<223> Accession number cg43975313

<400> 4879

gcacttagga tgttctggaa atgagaggaa atccacattc ctgccccagg a

51

<210> 4880

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (4879 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43975313

<400> 4880
gcacttagga tgttctggaa atgaggggaa atccacattc ctgccccagg a 51

<210> 4881
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4882 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43975716

<400> 4881
gaggatccag ttcaggttga tgtcctggga accagtgagg ctctcgctta a 51

<210> 4882
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4881 is other entry)

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<222> (0)...(0)
<223> Accession number cg43975716

<400> 4882
gaggatccag ttcaggttga tgtccaggga accagtgagg ctctcgctta a 51

<210> 4883
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4884 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43975716

<400> 4883
gttcaggttg atgtcctggg aaccagtgag gctctcgctt aaatctatgt g 51

<210> 4884
<211> 51
<212> DNA

<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4883 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43975716

<400> 4884
gttcagggtg atgtcctggg aaccaatgag gctctcgctt aaatctatgt g 51

<210> 4885
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4886 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43975716

<400> 4885
gtcctgggaa ccagtgaggc tctcgcttaa atctatgtgt tgaaagtggc a 51

<210> 4886
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4885 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43975716

<400> 4886
gtcctgggaa ccagtgaggc tctcgcttaa atctatgtgt tgaaagtggc a 51

<210> 4887
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4888 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43975716

<400> 4887
ggaaccagtg aggctctcgc ttaaactctat gtgttgaaag tggcattttg t 51

<210> 4888
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4887 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43975716

<400> 4888
ggaaccagtg aggctctcgc ttaaacctat gtgttgaaag tggcattttg t 51

<210> 4889
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4890 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43975716

<400> 4889
gctctcgctt aaatctatgt gttgaaagtg gcattttgtt gttgtagctt g 51

<210> 4890
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4889 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43975716

<400> 4890

gctctcgctt aaatctatgt gttgatagtg gcattttggtt gttgtagctt g

51

<210> 4891

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (4892 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43975716

<400> 4891

ctctcgctta aatctatgtg ttgaaagtggt cattttgttg ttgtagcttg t

51

<210> 4892

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (4891 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43975716

<400> 4892

ctctcgctta aatctatgtg ttgaatgtgg cattttgttg ttgtagcttg t

51

<210> 4893

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (4894 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43975716

<400> 4893

cttaaactta tgtgttgaaa gtggcatttt gttgtttag cttgtatcat a

51

<210> 4894

<211> 51

<212> DNA

<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4893 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43975716

<400> 4894
cttaaactcta tgtgttgaaa gtggcttttt gttgttgtag cttgtatcat a 51

<210> 4895
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4896 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43975716

<400> 4895
tctatgtggtt gaaagtggca ttttgttggt gtagcttgta tcatatgttg t 51

<210> 4896
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4895 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43975716

<400> 4896
tctatgtggtt gaaagtggca ttttgttggt gtagcttgta tcatatgttg t 51

<210> 4897
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4898 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43975716

<400> 4897
agtggcattt tgttggtgta gcttgatca tatgtgttt caaagtagtg t 51

<210> 4898
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4897 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43975716

<400> 4898
agtggcattt tgttggtgta gcttgatca tatgtgttt caaagtagtg t 51

<210> 4899
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4900 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43975716

<400> 4899
gcattttggt gttgtagctt gtatcatatg ttgtttcaaa gtagtgtcca a 51

<210> 4900
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4899 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43975716

<400> 4900
gcattttggt gttgtagctt gtatcgatatg ttgtttcaaa gtagtgtcca a 51

<210> 4901
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4902 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43975716

<400> 4901
tgtagcttgt atcatatggt gtttcaaagt agtgtccaaa ctgggaaggc a 51

<210> 4902
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4901 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43975716

<400> 4902
tgtagcttgt atcatatggt gtttcgaagt agtgtccaaa ctgggaaggc a 51

<210> 4903
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4904 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43975716

<400> 4903
gcttgatca tatgttgttt caaagtagtg tccaaactgg gaaggcaagg g 51

<210> 4904
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4903 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43975716

<400> 4904
gcttgatca tatgttgttt caaagcagtg tccaaactgg gaaggcaagg g 51

<210> 4905
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4906 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43975716

<400> 4905
gggaaggcaa gggtttgggc tgcacgatat tcctcaggtc atatctttcc t 51

<210> 4906
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4905 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43975716

<400> 4906
gggaaggcaa gggtttgggc tgcacaatat tcctcaggtc atatctttcc t 51

<210> 4907
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4908 is other entry)

<221> misc_feature

<222> (0)...(0)
<223> Accession number cg43975716

<400> 4907
gtttgggctg cacgatattc ctcagggtcat atctttcctg gttccagttg c 51

<210> 4908
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4907 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43975716

<400> 4908
gtttgggctg cacgatattc ctcagatcat atctttcctg gttccagttg c 51

<210> 4909
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4910 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43975716

<400> 4909
tgcccatcag ggtacggttt gaataggtat tctcattagt ggtgcatcgc c 51

<210> 4910
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4909 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43975716

<400> 4910
tgcccatcag ggtacggttt gaataggtat tctcattagt ggtgcatcgc c 51

<210> 4911
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4912 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43975856

<400> 4911
tggtgctaatt gtgctcaact ccttcacggg ggagagatca gtttgcaaag t 51

<210> 4912
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4911 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43975856

<400> 4912
tggtgctaatt gtgctcaact ccttcccggg ggagagatca gtttgcaaag t 51

<210> 4913
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4914 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43975884

<400> 4913
aaggcatata cacacctcat cccccacat gcacatcagc aagtctatca g 51

<210> 4914
<211> 51
<212> DNA
<213> Homo sapiens

<220>

<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4913 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43975884

<400> 4914
aaggcatata cacacctcat cccctacat gcacatcagc aagtctatca g 51

<210> 4915
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4916 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43975884

<400> 4915
tcattgggcc aaatgtttgg catatcagaa ttgtgatgt gagagggcaa g 51

<210> 4916
<211> 50
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4915 is other entry)

<221> misc_feature
<222> (25)...(26)
<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43975884

<400> 4916
tcattgggcc aaatgtttgg catatagaat ttgtgatgtg agagggcaag 50

<210> 4917
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)

<223> 1 of 2 allelic variants (4918 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43976147

<400> 4917

cctctctctct gtctctctct cccatgtgct gcaaactcac ccctgatgt c

51

<210> 4918

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (4917 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43976147

<400> 4918

cctctctctct gtctctctct cccatgtgct gcaaactcac ccctgatgt c

51

<210> 4919

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (4920 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43976147

<400> 4919

caaactcacc ccctgatgtc ccagcagcca gccctgggc ccaggttggt c

51

<210> 4920

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (4919 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43976147

<400> 4920
caaactcacc ccctgatgtc ccagctgcca gccctgggc ccaggttggt c 51

<210> 4921
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4922 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43976147

<400> 4921
aaactcacc cctgatgtcc cagcagccag ccctgggcc caggttggtc g 51

<210> 4922
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4921 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43976147

<400> 4922
aaactcacc cctgatgtcc cagcaccag ccctgggcc caggttggtc g 51

<210> 4923
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4924 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43976147

<400> 4923
gatgtcccag cagccagccc ctgggccag gttggtcgtt tctcctgctt t 51

<210> 4924
<211> 51
<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (4923 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43976147

<400> 4924

gatgtcccag cagccagccc ctggggccag gttggtcgtt tctcctgctt t

51

<210> 4925

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (4926 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43976147

<400> 4925

agcacggtac tgtcccacct gcaccatgct gcattctctc agcaccacct g

51

<210> 4926

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (4925 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43976147

<400> 4926

agcacggtac tgtcccacct gcaccctgct gcattctctc agcaccacct g

51

<210> 4927

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (4928 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43976147

<400> 4927
accatgctgc attctctcag caccacctgt gattctgccca ctggcctaata t 51

<210> 4928
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4927 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43976147

<400> 4928
accatgctgc attctctcag caccatctgt gattctgccca ctggcctaata t 51

<210> 4929
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4930 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43976147

<400> 4929
ttctctcagc accacctgtg attctgccac tggcctaatt cgtgatctca g 51

<210> 4930
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4929 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43976147

<400> 4930

ttctctcagc accacctgtg attctccac tggcctaatt cgtgatctca g

51

<210> 4931

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (4932 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43976643

<400> 4931

tggctgtcc agaagaacta catctttatt tagcatgcag tttgtggcaa t

51

<210> 4932

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (4931 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43976643

<400> 4932

tggctgtcc agaagaacta catctatatt tagcatgcag tttgtggcaa t

51

<210> 4933

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (4934 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43976643

<400> 4933

ggtctgtcca gaagaactac atctttatatt agcatgcagt ttgtggcaat t

51

<210> 4934

<211> 51

<212> DNA

<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4933 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43976643

<400> 4934
ggctctgtcca gaagaactac atcttaattt agcatgcagt ttgtggcaat t 51

<210> 4935
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4936 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43976643

<400> 4935
agtttgtggc aattctattg aggaggaaac aggatagtat gtacagtaac a 51

<210> 4936
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4935 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43976643

<400> 4936
agtttgtggc aattctattg aggagcaaac aggatagtat gtacagtaac a 51

<210> 4937
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4938 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43976643

<400> 4937
atgcagattc ttggcattcc attccgtttc tgctgattat ttcagaatta t 51

<210> 4938
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4937 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43976643

<400> 4938
atgcagattc ttggcattcc attccatttc tgctgattat ttcagaatta t 51

<210> 4939
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4940 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43976643

<400> 4939
ttcttggcat tccattccgt ttctgctgat tatttcagaa ttatctgttg g 51

<210> 4940
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (4939 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43976643

<400> 4940
ttcttggcat tccattccgt ttctgttgat tatttcagaa ttatctgttg g 51

<210> 4941
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4942 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43976643

<400> 4941
tgctgattat ttcagaatta tctgttgga tgaacgcggc gtgttgctgc a 51

<210> 4942
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<223> Accession number cg43976643

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<210> 4943
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<223> Accession number cg43976643

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cagaattatc tggtggaatg aacgcggcgt gttgctgcaa ctgtgcgctg g 51

<210> 4944
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<221> misc_feature
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<400> 4944
cagaattatc tgttggaatg aacgccgcgt gttgctgcaa ctgtgcgctg g 51

<210> 4945
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<223> 1 of 2 allelic variants (4946 is other entry)

<221> misc_feature
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<400> 4945
tatctgttgg aatgaacgcg gcgtgttgct gcaactgtgc gctggtgaga g 51

<210> 4946
<211> 51
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<213> Homo sapiens

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<223> 2 of 2 allelic variants (4945 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43976643

<400> 4946
tatctgttgg aatgaacgcg gcgtgctgct gcaactgtgc gctggtgaga g 51

<210> 4947
<211> 51
<212> DNA
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<223> 1 of 2 allelic variants (4948 is other entry)

<221> misc_feature

<222> (0)...(0)
<223> Accession number cg43976643

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ggaatgaacg cggcgtgttg ctgcaactgt gcgctggtga gagcctgctg g 51

<210> 4948
<211> 51
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<213> Homo sapiens

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<223> 2 of 2 allelic variants (4947 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43976643

<400> 4948
ggaatgaacg cggcgtgttg ctgcagctgt gcgctggtga gagcctgctg g 51

<210> 4949
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 1 of 2 allelic variants (4950 is other entry)

<221> misc_feature
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<223> Accession number cg43976643

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tgaacgcggc gtgttgctgc aactgtgcgc tggtagagagc ctgctggtag t 51

<210> 4950
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 2 of 2 allelic variants (4949 is other entry)

<221> misc_feature
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tgaacgcggc gtgttgctgc aactgcgcgc tggtagagagc ctgctggtag t 51

<210> 4951
<211> 51
<212> DNA
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<221> misc_feature
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<210> 4952
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 2 of 2 allelic variants (4951 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43976643

<400> 4952
cgtgttgctg caactgtgcg ctggtcagag cctgctggta gtgcaagacg c 51

<210> 4953
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 1 of 2 allelic variants (4954 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43976643

<400> 4953
gctggtagtg caagacgctg gggttaaaga ccgcgctggg accattgctt t 51

<210> 4954
<211> 51
<212> DNA
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<223> 2 of 2 allelic variants (4953 is other entry)

<221> misc_feature
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<223> Accession number cg43976643

<400> 4954
gctggtagtgc caagacgctg gggttgaaga cgcgctggt accattgctt t 51

<210> 4955
<211> 51
<212> DNA
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<223> 1 of 2 allelic variants (4956 is other entry)

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<223> Accession number cg43976643

<400> 4955
tttcaagtgc ttgtctcttt ggtaaaggat gaagagcacc aggggggaaag g 51

<210> 4956
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (4955 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43976643

<400> 4956
tttcaagtgc ttgtctcttt ggtaagggat gaagagcacc aggggggaaag g 51

<210> 4957
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 1 of 2 allelic variants (4958 is other entry)

<221> misc_feature
<222> (0)...(0)

<223> Accession number cg43976707

<400> 4957

gaatgcgttt ccccttcag gaagagaact cagttacaca tcacgaagtc a

51

<210> 4958

<211> 50

<212> DNA

<213> Homo sapiens

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<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (4957 is other entry)

<221> misc_feature

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<222> (0)...(0)

<223> Accession number cg43976707

<400> 4958

gaatgcgttt ccccttcag gaagaaactc agttacacat cacgaagtca

50

<210> 4959

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 1 of 2 allelic variants (4960 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43976754

<400> 4959

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<210> 4960

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 2 of 2 allelic variants (4959 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43976754

<400> 4960
gatgcaaagc gacctggaca aggcttacct cagtgccagg agggtcacct g 51

<210> 4961
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<212> DNA
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<223> 1 of 2 allelic variants (4962 is other entry)

<221> misc_feature
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<223> Accession number cg43976910

<400> 4961
ctgggtgttca tcgtggctga aggggactcc ctgggtgtgct acaatccctt g 51

<210> 4962
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (4961 is other entry)

<221> misc_feature
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<223> Accession number cg43976910

<400> 4962
ctgggtgttca tcgtggctga agggggctcc ctgggtgtgct acaatccctt g 51

<210> 4963
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4964 is other entry)

<221> misc_feature
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<223> Accession number cg43976910

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actgctgctg ctcagctggt tgcttgaact gacagtaggc cagcctgttc t 51

<210> 4964
<211> 51
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<213> Homo sapiens

<220>

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<223> 2 of 2 allelic variants (4963 is other entry)

<221> misc_feature

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<223> Accession number cg43976910

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actgctgctg ctcagctggt tgcttaaact gacagtaggc cagcctgttc t

51

<210> 4965

<211> 50

<212> DNA

<213> Homo sapiens

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<222> (26)...(0)

<223> 1 of 2 allelic variants (4966 is other entry)

<221> misc_feature

<222> (25)...(26)

<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43977247

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ttggcctcct tgctgtcctg cttcttggtt tcagaaactt ctgacttccg

50

<210> 4966

<211> 51

<212> DNA

<213> Homo sapiens

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<222> (26)...(0)

<223> 2 of 2 allelic variants (4965 is other entry)

<221> misc_feature

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<223> Accession number cg43977247

<400> 4966

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51

<210> 4967

<211> 51

<212> DNA

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<223> 1 of 2 allelic variants (4968 is other entry)

<221> misc_feature
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<223> Accession number cg43977442

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ttctgcctgt gctcctcaag ggcagctggt gaggcctgga attcctcact g 51

<210> 4968
<211> 51
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<223> 2 of 2 allelic variants (4967 is other entry)

<221> misc_feature
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<223> Accession number cg43977442

<400> 4968
ttctgcctgt gctcctcaag ggcagatggt gaggcctgga attcctcact g 51

<210> 4969
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (4970 is other entry)

<221> misc_feature
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<223> Accession number cg43977442

<400> 4969
agtcccgggc tcgattcaca ggctcatgct ccaggtgctg caggctgttg g 51

<210> 4970
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4969 is other entry)

<221> misc_feature

<222> (0)...(0)
<223> Accession number cg43977442

<400> 4970
agtcccgggc tcgattcaca ggctcgtgct ccagggtgctg caggctgttg g 51

<210> 4971
<211> 51
<212> DNA
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<223> 1 of 2 allelic variants (4972 is other entry)

<221> misc_feature
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<223> Accession number cg43979051

<400> 4971
taaagataaa tgaagaattc aaaaataata aaagtgaac ttcttctaag a 51

<210> 4972
<211> 51
<212> DNA
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<223> 2 of 2 allelic variants (4971 is other entry)

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<223> Accession number cg43979051

<400> 4972
taaagataaa tgaagaattc aaaaaaata aaagtgaac ttcttctaag a 51

<210> 4973
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 1 of 2 allelic variants (4974 is other entry)

<221> misc_feature
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<223> Accession number cg43979107

<400> 4973
gtaggccaga atataaatct ccttaggaag aagtttggtt aggcaaaacc a 51

<210> 4974
<211> 50
<212> DNA
<213> Homo sapiens

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<223> 2 of 2 allelic variants (4973 is other entry)

<221> misc_feature
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<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43979107

<400> 4974
gtaggccaga atataaatct ccttagaaga agtttggtta ggcaaaacca

50

<210> 4975
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4976 is other entry)

<221> misc_feature
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<400> 4975
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51

<210> 4976
<211> 50
<212> DNA
<213> Homo sapiens

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<223> 2 of 2 allelic variants (4975 is other entry)

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<223> Nucleotide deleted between bases 25 and 26

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<222> (0)...(0)
<223> Accession number cg43979107

<400> 4976

taggccagaa tataaatctc cttagaagaa gtttggttag gcaaaaccac

50

<210> 4977

<211> 51

<212> DNA

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<223> 1 of 2 allelic variants (4978 is other entry)

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51

<210> 4978

<211> 51

<212> DNA

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<223> 2 of 2 allelic variants (4977 is other entry)

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<223> Accession number cg43979152

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ttctttggac cttttccaga acatgggtact tccctgcctc agttagaagt t

51

<210> 4979

<211> 51

<212> DNA

<213> Homo sapiens

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<222> (26)...(0)

<223> 1 of 2 allelic variants (4980 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43979411

<400> 4979

gcacagtggc tcatgcctgt aatcctagca ctttgggagg ctgaggcagg t

51

<210> 4980

<211> 51

<212> DNA

<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4979 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43979411

<400> 4980
gcacagtggc tcatgcctgt aatcccagca ctttgggagg ctgaggcagg t 51

<210> 4981
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4982 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43979439

<400> 4981
gaaatcagaa agcaaggctt gttttaggca aacgtgccac atatcttgta a 51

<210> 4982
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4981 is other entry)

<221> misc_feature
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<223> Accession number cg43979439

<400> 4982
gaaatcagaa agcaaggctt gttttgggca aacgtgccac atatcttgta a 51

<210> 4983
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4984 is other entry)

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<222> (0)...(0)
<223> Accession number cg43979495

<400> 4983
agagtgcacg tctaaagggtg gttcacaagg gaaggatcca caagttagtg a 51

<210> 4984
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (4983 is other entry)

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<400> 4984
agagtgcacg tctaaagggtg gttcagaagg gaaggatcca caagttagtg a 51

<210> 4985
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (4986 is other entry)

<221> misc_feature
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<223> Accession number cg43979568

<400> 4985
agcgggtgagg ggcgctcgaa gaagggcctc tgggccagcg gggagcgag c 51

<210> 4986
<211> 50
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (4985 is other entry)

<221> misc_feature
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<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)

<223> Accession number cg43979568

<400> 4986

agcggtagg ggcctcgaa gaaggcctct gggccagcgg ggagcgcagc

50

<210> 4987

<211> 51

<212> DNA

<213> Homo sapiens

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<222> (26)...(0)

<223> 1 of 2 allelic variants (4988 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43979574

<400> 4987

attgcaattc taatcaatca atcaaacag accaaaaggt catacttcta a

51

<210> 4988

<211> 50

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (4987 is other entry)

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<223> Nucleotide deleted between bases 25 and 26

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<222> (0)...(0)

<223> Accession number cg43979574

<400> 4988

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50

<210> 4989

<211> 51

<212> DNA

<213> Homo sapiens

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<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (4990 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43979594

<400> 4989
taaggccatt tcactaagtt tgggaattgga ggagaggcaa gccaagaatc a 51

<210> 4990
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (4989 is other entry)

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<223> Accession number cg43979594

<400> 4990
taaggccatt tcactaagtt tggaaatgga ggagaggcaa gccaagaatc a 51

<210> 4991
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (4992 is other entry)

<221> misc_feature
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<223> Accession number cg43979615

<400> 4991
ctgcagacca ctctgtggca cgggatgagg atggggacag gattgcctct c 51

<210> 4992
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (4991 is other entry)

<221> misc_feature
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<223> Accession number cg43979615

<400> 4992
ctgcagacca ctctgtggca cgggacgagg atggggacag gattgcctct c 51

<210> 4993
<211> 51
<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (4994 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43979647

<400> 4993

ttgctcagag cacaaggcct accccatcct gcgtctccag cccgacttgg a

51

<210> 4994

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (4993 is other entry)

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<223> Accession number cg43979647

<400> 4994

ttgctcagag cacaaggcct accccgtcct gcgtctccag cccgacttgg a

51

<210> 4995

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (4996 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43979647

<400> 4995

tctttattta caaaatgcgg tgaagagaaa atatctagat atttgggtgt a

51

<210> 4996

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (4995 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43979647

<400> 4996
tctttatttta caaaatgcgg tgaagggaaa atatctagat atttgggtgt a 51

<210> 4997
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (4998 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43979837

<400> 4997
gtggagtttg gggagggggg cgaaagcaac gggactgctg ggagaggagg g 51

<210> 4998
<211> 50
<212> DNA
<213> Homo sapiens

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<222> (26)...(0)
<223> 2 of 2 allelic variants (4997 is other entry)

<221> misc_feature
<222> (25)...(26)
<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43979837

<400> 4998
gtggagtttg gggagggggg cgaaacaacg ggactgctgg gagaggaggg 50

<210> 4999
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5000 is other entry)

<221> misc_feature

<222> (0)...(0)
<223> Accession number cg43979896

<400> 4999
taagaccaat tcagaacaaa ggcaggttgc ccttaaaaca ggtttgacct t 51

<210> 5000
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (4999 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43979896

<400> 5000
taagaccaat tcagaacaaa ggcagattgc ccttaaaaca ggtttgacct t 51

<210> 5001
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5002 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43980073

<400> 5001
atttatgaag aaatggactt ggaaaggaaa ttctaacaga gaagagctta a 51

<210> 5002
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 2 of 2 allelic variants (5001 is other entry)

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<223> Accession number cg43980073

<400> 5002
atttatgaag aaatggactt ggaaatgaaa ttctaacaga gaagagctta a 51

<210> 5003
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 1 of 2 allelic variants (5004 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43980088

<400> 5003
aatcacatt tagtattata gaaaaagagt ttctcagtaa tctcttgacc a 51

<210> 5004
<211> 50
<212> DNA
<213> Homo sapiens

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<223> 2 of 2 allelic variants (5003 is other entry)

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<222> (25)...(26)
<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43980088

<400> 5004
aatcacatt tagtattata gaaaagagtt tctcagtaat ctcttgacca 50

<210> 5005
<211> 50
<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (5006 is other entry)

<221> misc_feature
<222> (25)...(26)
<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43980276

<400> 5005

tttcaaatat aaatcattta actataaata ttcagaggac attcaggaga

50

<210> 5006

<211> 51

<212> DNA

<213> Homo sapiens

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<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (5005 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43980276

<400> 5006

tttcaaatat aaatcattta actataaaat attcagagga cattcaggag a

51

<210> 5007

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

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<223> 1 of 2 allelic variants (5008 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43980513

<400> 5007

aacgcatgcc gcctcggccg tcagggcgct gatctcgccg ttgaggggtgc t

51

<210> 5008

<211> 50

<212> DNA

<213> Homo sapiens

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<222> (26)...(0)

<223> 2 of 2 allelic variants (5007 is other entry)

<221> misc_feature

<222> (25)...(26)

<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43980513

<400> 5008

aacgcatgcc gcctcggccg tcagggcgctg atctcgccgt tgaggggtgct

50

<210> 5009
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (5010 is other entry)

<221> misc_feature
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<223> Accession number cg43980531

<400> 5009
tcttcctctg ttgtcatcgg tgaagctaaa aaaaagtttt ctgaaagtag c 51

<210> 5010
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5009 is other entry)

<221> misc_feature
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<223> Accession number cg43980531

<400> 5010
tcttcctctg ttgtcatcgg tgaagataaa aaaaagtttt ctgaaagtag c 51

<210> 5011
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 1 of 2 allelic variants (5012 is other entry)

<221> misc_feature
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<223> Accession number cg43980531

<400> 5011
gttgcacatg gtgaagctaa aaaaagtttt tctgaaagta gcaagttgtg t 51

<210> 5012
<211> 50
<212> DNA
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<220>

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<222> (26)...(0)
<223> 2 of 2 allelic variants (5011 is other entry)

<221> misc_feature
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<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43980531

<400> 5012
gttgtcatcg gtgaagctaa aaaaagtttt ctgaaagtag caagttgtgt 50

<210> 5013
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5014 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43980531

<400> 5013
gccaaaaagg ctcagtcttt ggctcacaga tgtcagtgac aaaatcatgg c 51

<210> 5014
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5013 is other entry)

<221> misc_feature
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<223> Accession number cg43980531

<400> 5014
gccaaaaagg ctcagtcttt ggctcgacaga tgtcagtgac aaaatcatgg c 51

<210> 5015
<211> 51
<212> DNA
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<222> (26)...(0)

<223> 1 of 2 allelic variants (5016 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43980543

<400> 5015

attattgaat actttttgag tatttgctat ataccaggca aaaggcacag a

51

<210> 5016

<211> 51

<212> DNA

<213> Homo sapiens

<220>

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<222> (26)...(0)

<223> 2 of 2 allelic variants (5015 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43980543

<400> 5016

attattgaat actttttgag tattttactat ataccaggca aaaggcacag a

51

<210> 5017

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (5018 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43980543

<400> 5017

ctttttgagt atttgctata taccaggcaa aaggcacaga acaaattatt t

51

<210> 5018

<211> 51

<212> DNA

<213> Homo sapiens

<220>

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<222> (26)...(0)

<223> 2 of 2 allelic variants (5017 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43980543

<400> 5018
cttttttgagt atttgctata taccaagcaa aaggcacaga acaaattatt t 51

<210> 5019
<211> 51
<212> DNA
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5020 is other entry)

<221> misc_feature
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<223> Accession number cg43980655

<400> 5019
cacccaggct ggagtgcagt ggcgcgatct cggctcactg caacctccgc c 51

<210> 5020
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<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5019 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43980655

<400> 5020
cacccaggct ggagtgcagt ggcgcaatct cggctcactg caacctccgc c 51

<210> 5021
<211> 50
<212> DNA
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5022 is other entry)

<221> misc_feature
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<221> misc_feature
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<223> Accession number cg43980827

<400> 5021
ttcatatcaa aattcccata aaaaattaca ttccccccctc cccagttcta 50

<210> 5022

<211> 51

<212> DNA

<213> Homo sapiens

<220>

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<222> (26)...(0)

<223> 2 of 2 allelic variants (5021 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43980827

<400> 5022

ttcatatcaa aattcccata aaaaaattac attccccct cccagttct a

51

<210> 5023

<211> 51

<212> DNA

<213> Homo sapiens

<220>

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<222> (26)...(0)

<223> 1 of 2 allelic variants (5024 is other entry)

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<223> Accession number cg43980859

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actggttagga agagcagaga gatgcaggga acgtggtcag aggctgtgaa c

51

<210> 5024

<211> 51

<212> DNA

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<223> 2 of 2 allelic variants (5023 is other entry)

<221> misc_feature

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<223> Accession number cg43980859

<400> 5024

actggttagga agagcagaga gatgcgggga acgtggtcag aggctgtgaa c

51

<210> 5025

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 1 of 2 allelic variants (5026 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43981033

<400> 5025
gattgttggt aacactgaaa aaaaacatg gtggctcctg aaacaagaca g 51

<210> 5026
<211> 50
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5025 is other entry)

<221> misc_feature
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<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43981033

<400> 5026
gattgttggt aacactgaaa aaaaacatgg tggctcctga aacaagacag 50

<210> 5027
<211> 51
<212> DNA
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5028 is other entry)

<221> misc_feature
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<223> Accession number cg43981033

<400> 5027
aaaacatggt ggctcctgaa acaagacagg ttagcaactg gtacagcttt c 51

<210> 5028
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5027 is other entry)

<221> misc_feature
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<223> Accession number cg43981033

<400> 5028
aaaacatggt ggctcctgaa acaaggcagg ttagcaactg gtacagcttt c 51

<210> 5029
<211> 51
<212> DNA
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<223> 1 of 2 allelic variants (5030 is other entry)

<221> misc_feature
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<223> Accession number cg43981094

<400> 5029
ttcataagac ttgagtaagt agatccaaat ttgttatcac taatggctca a 51

<210> 5030
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (5029 is other entry)

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<223> Accession number cg43981094

<400> 5030
ttcataagac ttgagtaagt agatccaaat ttgttatcac taatggctca a 51

<210> 5031
<211> 51
<212> DNA
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5032 is other entry)

<221> misc_feature
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<223> Accession number cg43981141

<400> 5031
gaacacggtg ctagcagcta acgaccttcc catgtgccct gtcctcctga c 51

<210> 5032
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (5031 is other entry)

<221> misc_feature
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<223> Accession number cg43981141

<400> 5032
gaacacggtg ctagcagcta acgactttcc catgtgccct gtcctcctga c 51

<210> 5033
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<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5034 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43981141

<400> 5033
ggagggtgga actatgtttg aaggccttcc cacactctgt acattcataa g 51

<210> 5034
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (5033 is other entry)

<221> misc_feature
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<400> 5034
ggagggtgga actatgtttg aaggctttcc cacactctgt acattcataa g 51

<210> 5035
<211> 51

<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5036 is other entry)

<221> misc_feature
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<223> Accession number cg43981267

<400> 5035
gccaggagga ccagcatggg gacacaggag ctatcaggca ggagtcaggc t 51

<210> 5036
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5035 is other entry)

<221> misc_feature
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<400> 5036
gccaggagga ccagcatggg gacaccggag ctatcaggca ggagtcaggc t 51

<210> 5037
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5038 is other entry)

<221> misc_feature
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<223> Accession number cg43981280

<400> 5037
attttgttct gcaagataat ttacatagtt ctctctgcta tatgtgtcca c 51

<210> 5038
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (5037 is other entry)

<221> misc_feature

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<223> Accession number cg43981280

<400> 5038

attttgttct gcaagataat ttacacagtt ctctctgcta tatgtgtcca c

51

<210> 5039

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 1 of 2 allelic variants (5040 is other entry)

<221> misc_feature

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<223> Accession number cg43981323

<400> 5039

gtggctcccc tggggacagg gcctctgcg gcgcggcat cccctctcg g

51

<210> 5040

<211> 50

<212> DNA

<213> Homo sapiens

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<222> (26)...(0)

<223> 2 of 2 allelic variants (5039 is other entry)

<221> misc_feature

<222> (25)...(26)

<223> Nucleotide deleted between bases 25 and 26

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<222> (0)...(0)

<223> Accession number cg43981323

<400> 5040

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<210> 5041

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (5042 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43981356

<400> 5041
tgctgctgcc acggggtagg ggtgcgggag gcggcctggc ctcatggccg c 51

<210> 5042
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5041 is other entry)

<221> misc_feature
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<223> Accession number cg43981356

<400> 5042
tgctgctgcc acggggtagg ggtgcaggag gcggcctggc ctcatggccg c 51

<210> 5043
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5044 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43981449

<400> 5043
tccaatggga tcttagaatt cctagagttg cagaatttta ctgactagac c 51

<210> 5044
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5043 is other entry)

<221> misc_feature
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<223> Accession number cg43981449

<400> 5044
tccaatggga tcttagaatt cctagcgttg cagaatttta ctgactagac c 51

<210> 5045
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5046 is other entry)

<221> misc_feature
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<400> 5045
caaggagct cataactcca tggggccag agttcctggc tcgggaacat t 51

<210> 5046
<211> 51
<212> DNA
<213> Homo sapiens

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<222> (26)...(0)
<223> 2 of 2 allelic variants (5045 is other entry)

<221> misc_feature
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<223> Accession number cg43981449

<400> 5046
caaggagct cataactcca tggggccag agttcctggc tcgggaacat t 51

<210> 5047
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5048 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43981504

<400> 5047
ttttaacccc ccaaacttta aattcaatat tatatattct atttgaattt t 51

<210> 5048
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5047 is other entry)

<221> misc_feature
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<223> Accession number cg43981504

<400> 5048
ttttaacccc ccaaattctta aattcgatat tatatatattct atttgaattt t 51

<210> 5049
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5050 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43981615

<400> 5049
aatcttgagg gctgaagggt ccaaagaaat ggtatatata gaattctatc t 51

<210> 5050
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 2 of 2 allelic variants (5049 is other entry)

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aatcttgagg gctgaagggt ccaaataaat ggtatatata gaattctatc t 51

<210> 5051
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<223> 1 of 2 allelic variants (5052 is other entry)

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<210> 5052
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<212> DNA
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<220>
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<223> 2 of 2 allelic variants (5051 is other entry)

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<400> 5052
aaagaaatgg tatatataga attctctctg acttgaaatt ttcccttct g 51

<210> 5053
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5054 is other entry)

<221> misc_feature
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<223> Accession number cg43981615

<400> 5053
ggtatatata gaattctatc tgacttgaaa tttcccttc ctggagctcc g 51

<210> 5054
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5053 is other entry)

<221> misc_feature
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<223> Accession number cg43981615

<400> 5054
ggtatatata gaattctatc tgactcgaaa tttcccttc ctggagctcc g 51

<210> 5055
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (5056 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43981615

<400> 5055
atagaattct atctgacttg aaattttccc ttcttgaggc tccggatgct g 51

<210> 5056
<211> 51
<212> DNA
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<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5055 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43981615

<400> 5056
atagaattct atctgacttg aaattgtccc ttcttgaggc tccggatgct g 51

<210> 5057
<211> 51
<212> DNA
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<223> 1 of 2 allelic variants (5058 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43981615

<400> 5057
actgagggtca tgatgttgga atcttgaggg ctgaagggttc caaagaaatg g 51

<210> 5058
<211> 51
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<223> 2 of 2 allelic variants (5057 is other entry)

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<223> Accession number cg43981615

<400> 5058
actgaggtca tgatgttgga atcttaaggg ctgaagggttc caaagaaatg g 51

<210> 5059
<211> 51
<212> DNA
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<223> 1 of 2 allelic variants (5060 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43981624

<400> 5059
tggcagcccc tgcttttttt ttttttttag ctccctaaag actgtagcag g 51

<210> 5060
<211> 50
<212> DNA
<213> Homo sapiens

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<223> 2 of 2 allelic variants (5059 is other entry)

<221> misc_feature
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<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43981624

<400> 5060
tggcagcccc tgcttttttt ttttttttagc tccctaaaga ctgtagcagg 50

<210> 5061
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (5062 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43981624

<400> 5061

ggcagccctt gctttttttt ttttttagc tcctaaaga ctgtagcagg a

51

<210> 5062

<211> 50

<212> DNA

<213> Homo sapiens

<220>

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<223> 2 of 2 allelic variants (5061 is other entry)

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<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43981624

<400> 5062

ggcagccctt gctttttttt ttttttagct ccctaaagac tgtagcagga

50

<210> 5063

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 1 of 2 allelic variants (5064 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43981624

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51

<210> 5064

<211> 50

<212> DNA

<213> Homo sapiens

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<223> 2 of 2 allelic variants (5063 is other entry)

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<223> Accession number cg43981624

<400> 5064
gcagcccctg cttttttttt ttttttagctc cctaaagact gtagcaggat 50

<210> 5065
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (5066 is other entry)

<221> misc_feature
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<223> Accession number cg43981644

<400> 5065
ctgtttgctt gttggtgtga gtttttcttc tggagacttt gtactgaatg t 51

<210> 5066
<211> 50
<212> DNA
<213> Homo sapiens

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<223> 2 of 2 allelic variants (5065 is other entry)

<221> misc_feature
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<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43981644

<400> 5066
ctgtttgctt gttggtgtga gtttttcttc ggagactttg tactgaatgt 50

<210> 5067
<211> 49
<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (5068 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43981789

<400> 5067
cggggagaac gacagttgct gcagggaatc ttttaaacga gagcgagaa 49

<210> 5068
<211> 49
<212> DNA
<213> Homo sapiens

<220>
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<222> (24)...(0)
<223> 2 of 2 allelic variants (5067 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43981789

<400> 5068
cggggagaac gacagttgct gcaaggaatc ttttaaacga gagcgagaa 49

<210> 5069
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5070 is other entry)

<221> misc_feature
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<223> Accession number cg43981889

<400> 5069
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<210> 5070
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5069 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43981889

<400> 5070
ctaattctggt gtgaaagatc ttttatttgt atttccattc ttcgattatc t 51

<210> 5071
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (5072 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43981889

<400> 5071
gattatctct ctcaagtaca gatgattgct tgttggcttt atcagtgtcc t 51

<210> 5072
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 2 of 2 allelic variants (5071 is other entry)

<221> misc_feature
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<223> Accession number cg43981889

<400> 5072
gattatctct ctcaagtaca gatgaatgct tgttggcttt atcagtgtcc t 51

<210> 5073
<211> 51
<212> DNA
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<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5074 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43981997

<400> 5073
agtagtactg gagggccttg aggggcccac agacagatcc catccatcag c 51

<210> 5074
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<212> DNA

<213> Homo sapiens

<220>

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<222> (26)...(0)

<223> 2 of 2 allelic variants (5073 is other entry)

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<222> (0)...(0)

<223> Accession number cg43981997

<400> 5074

agtagtactg gagggccttg agggggccaca gacagatccc atccatcagc

50

<210> 5075

<211> 51

<212> DNA

<213> Homo sapiens

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<222> (26)...(0)

<223> 1 of 2 allelic variants (5076 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43982038

<400> 5075

aaatgcttct ctctctttgg tctccctaag gtcctcctcc tagtacacag g

51

<210> 5076

<211> 50

<212> DNA

<213> Homo sapiens

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<222> (26)...(0)

<223> 2 of 2 allelic variants (5075 is other entry)

<221> misc_feature

<222> (25)...(26)

<223> Nucleotide deleted between bases 25 and 26

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<223> Accession number cg43982038

<400> 5076

aaatgcttct ctctctttgg tctcctaagg tctcctcct agtacacagg

50

<210> 5077

<211> 51
<212> DNA
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<220>
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<223> 1 of 2 allelic variants (5078 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43982123

<400> 5077
acacacacac acacacacac acacacacac acagcaccat gtcctgagct g 51

<210> 5078
<211> 50
<212> DNA
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<223> 2 of 2 allelic variants (5077 is other entry)

<221> misc_feature
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<223> Nucleotide deleted between bases 25 and 26

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<223> Accession number cg43982123

<400> 5078
acacacacac acacacacac acacaacaca cagcaccatg tcctgagctg 50

<210> 5079
<211> 51
<212> DNA
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<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5080 is other entry)

<221> misc_feature
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<223> Accession number cg43982123

<400> 5079
acacacacac acacacacac acacacacac agcaccatgt cctgagctgc t 51

<210> 5080
<211> 50
<212> DNA

<213> Homo sapiens

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<222> (26)...(0)

<223> 2 of 2 allelic variants (5079 is other entry)

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<222> (0)...(0)

<223> Accession number cg43982123

<400> 5080

acacacacac acacacacac acacaacaca gcaccatgct ctgagctgct

50

<210> 5081

<211> 51

<212> DNA

<213> Homo sapiens

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<221> misc_feature

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<223> 1 of 2 allelic variants (5082 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43982123

<400> 5081

gactcccgcc agactgcacc tgacagacaa agccaactag gagggagggg a

51

<210> 5082

<211> 51

<212> DNA

<213> Homo sapiens

<220>

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<222> (26)...(0)

<223> 2 of 2 allelic variants (5081 is other entry)

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<222> (0)...(0)

<223> Accession number cg43982123

<400> 5082

gactcccgcc agactgcacc tgacacacaa agccaactag gagggagggg a

51

<210> 5083

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 1 of 2 allelic variants (5084 is other entry)

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<223> Accession number cg43982153

<400> 5083
ttaagtactt tcagtgtcga aaaaaatgca atcactgtgt tgtatataat a 51

<210> 5084
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<212> DNA
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<220>
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<223> 2 of 2 allelic variants (5083 is other entry)

<221> misc_feature
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<222> (0)...(0)
<223> Accession number cg43982153

<400> 5084
ttaagtactt tcagtgtcga aaaaatgcaa tcactgtggt gtatataata 50

<210> 5085
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5086 is other entry)

<221> misc_feature
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<223> Accession number cg43982189

<400> 5085
gcagctttgc gggcccgacc ggctccgcgt ggcaggtgaa gtgcaccggt t 51

<210> 5086
<211> 51
<212> DNA
<213> Homo sapiens

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<222> (26)...(0)
<223> 2 of 2 allelic variants (5085 is other entry)

<221> misc_feature
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<223> Accession number cg43982189

<400> 5086
gcagctttgc gggcccgacc ggctctgcgt ggcagggtgaa gtgcaccggt t 51

<210> 5087
<211> 51
<212> DNA
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<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5088 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43982232

<400> 5087
ggcaaaacac aatacaatga aatggaaaaa taatgtttgt tacaggagtgc c 51

<210> 5088
<211> 50
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5087 is other entry)

<221> misc_feature
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<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43982232

<400> 5088
ggcaaaacac aatacaatga aatggaaaat aatgtttgtt acaggagtgc 50

<210> 5089
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5090 is other entry)

<221> misc_feature
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<223> Accession number cg43982403

<400> 5089
tgcctaaaat ttctttctgga ataacatcat ctaatacact atatacatat t 51

<210> 5090
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (5089 is other entry)

<221> misc_feature
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<223> Accession number cg43982403

<400> 5090
tgcctaaaat ttctttctgga ataacgtcat ctaatacact atatacatat t 51

<210> 5091
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5092 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43982403

<400> 5091
tttcttctgg aataacatca tctaatacac tatatacata ttgtaaaac t 51

<210> 5092
<211> 51
<212> DNA
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<220>
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<221> misc_feature
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<400> 5092

tttcttctgg aataacatca tctaacacac tatatacata tttgtaaaac t 51

<210> 5093
<211> 51
<212> DNA
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<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5094 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43982403

<400> 5093
tatatacata tttgtaaaac ttatcaaacg aggtagacat gagttccata c 51

<210> 5094
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (5093 is other entry)

<221> misc_feature
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<223> Accession number cg43982403

<400> 5094
tatatacata tttgtaaaac ttatcgaacg aggtagacat gagttccata c 51

<210> 5095
<211> 51
<212> DNA
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<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5096 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43982403

<400> 5095
acttatcaaa cgaggtagac atgagttcca tacagatcca acagtcaccc t 51

<210> 5096
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (5095 is other entry)

<221> misc_feature
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<400> 5096
acttatcaaa cgaggtagac atgagctcca tacagatcca acagtcaccc t 51

<210> 5097
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5098 is other entry)

<221> misc_feature
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<223> Accession number cg43982403

<400> 5097
caccctctct gaagagtgca ccataaaact gaacaatgta tgggcaatca c 51

<210> 5098
<211> 51
<212> DNA
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<220>
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<223> 2 of 2 allelic variants (5097 is other entry)

<221> misc_feature
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<400> 5098
caccctctct gaagagtgca ccatagaact gaacaatgta tgggcaatca c 51

<210> 5099
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5100 is other entry)

<221> misc_feature
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<223> Accession number cg43982551

<400> 5099
tccataaaca gtgtcacccg caaaaaaagc catcacagata tcatgacggt g 51

<210> 5100
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5099 is other entry)

<221> misc_feature
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<223> Accession number cg43982551

<400> 5100
tccataaaca gtgtcacccg caaaacaagc catcacagata tcatgacggt g 51

<210> 5101
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5102 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43982652

<400> 5101
aaaaaaaaat taggcaggca tgggtggtggg cgctgtagt cccagctact c 51

<210> 5102
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5101 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43982652

<400> 5102
aaaaaaaaat taggcaggca tgggtgatggg cgctgtagt cccagctact c 51

<210> 5103
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5104 is other entry)

<221> misc_feature
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<223> Accession number cg43982782

<400> 5103
agttcgcgga caccgccatt tacaacgacg aggcggaccc gctggtgggc t 51

<210> 5104
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5103 is other entry)

<221> misc_feature
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<223> Accession number cg43982782

<400> 5104
agttcgcgga caccgccatt tacaatgacg aggcggaccc gctggtgggc t 51

<210> 5105
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5106 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43983113

<400> 5105
ttgactcatg gaggtctgt aagagcccag gattgggggc ctgggtcagg g 51

<210> 5106
<211> 50
<212> DNA
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<220>
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<223> 2 of 2 allelic variants (5105 is other entry)

<221> misc_feature
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<221> misc_feature
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<223> Accession number cg43983113

<400> 5106
ttgactcatg gaggtcttgt aagagccagg attgggggcc tgggtcaggg 50

<210> 5107
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5108 is other entry)

<221> misc_feature
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<400> 5107
agcccagaag gagaccatt ttttttggcg gggggagctg agtcccagag g 51

<210> 5108
<211> 50
<212> DNA
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<220>
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<223> 2 of 2 allelic variants (5107 is other entry)

<221> misc_feature
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<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
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<223> Accession number cg43983113

<400> 5108
agcccagaag gagaccatt ttttttggcg ggggagctga gtcccagagg 50

<210> 5109
<211> 51
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<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (5110 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43983122

<400> 5109

tgcacctgaa ggtgcccccc atgtggccaa agcccccttg cctgaccagg c

51

<210> 5110

<211> 50

<212> DNA

<213> Homo sapiens

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<223> 2 of 2 allelic variants (5109 is other entry)

<221> misc_feature

<222> (25)...(26)

<223> Nucleotide deleted between bases 25 and 26

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<222> (0)...(0)

<223> Accession number cg43983122

<400> 5110

tgcacctgaa ggtgcccccc atgtgccaaa gcccccttgc ctgaccaggc

50

<210> 5111

<211> 51

<212> DNA

<213> Homo sapiens

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<222> (26)...(0)

<223> 1 of 2 allelic variants (5112 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43983294

<400> 5111

caggggagct ggcacattcc tcagattctg gcatgtcatc ctggaagtac t

51

<210> 5112

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 2 of 2 allelic variants (5111 is other entry)

<221> misc_feature
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<400> 5112
caggggcgct ggcacattcc tcagactctg gcatgtcatc ctggaagtac t 51

<210> 5113
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (5114 is other entry)

<221> misc_feature
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<400> 5113
cagcctggcg gtactgccac agacgcaggt tcccgcccc cgaactgctg a 51

<210> 5114
<211> 51
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<220>
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<223> 2 of 2 allelic variants (5113 is other entry)

<221> misc_feature
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<223> Accession number cg43983294

<400> 5114
cagcctggcg gtactgccac agacgtaggt tcccgcccc cgaactgctg a 51

<210> 5115
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (5116 is other entry)

<221> misc_feature

<222> (0)...(0)
<223> Accession number cg43983294

<400> 5115
ctggcggtac tgccacagac gcagggtccc gtcccacgaa ctgctgacaa t 51

<210> 5116
<211> 51
<212> DNA
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<223> 2 of 2 allelic variants (5115 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43983294

<400> 5116
ctggcggtac tgccacagac gcagggtccc gtcccacgaa ctgctgacaa t 51

<210> 5117
<211> 51
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<213> Homo sapiens

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<223> 1 of 2 allelic variants (5118 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43983294

<400> 5117
ggtactgccca cagacgcagg ttcccgctccc acgaactgct gacaatcttt t 51

<210> 5118
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 2 of 2 allelic variants (5117 is other entry)

<221> misc_feature
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<223> Accession number cg43983294

<400> 5118
ggtactgccca cagacgcagg ttcccatccc acgaactgct gacaatcttt t 51

<210> 5119
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (5120 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43983486

<400> 5119
ctagcacacc ctctccaagg gtaccgcgctc gatgctatgt gctcagttct a 51

<210> 5120
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (5119 is other entry)

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<223> Accession number cg43983486

<400> 5120
ctagcacacc ctctccaagg gtaccacgctc gatgctatgt gctcagttct a 51

<210> 5121
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 1 of 2 allelic variants (5122 is other entry)

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<223> Accession number cg43983486

<400> 5121
aatgggatttt gtaaagtttt ttttttaatt tattcaaaaa aagacatagt a 51

<210> 5122
<211> 50
<212> DNA
<213> Homo sapiens

<220>

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<221> misc_feature
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<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43983486

<400> 5122
aatgggtattt gtaaagtttt tttttaattt attcaaaaaa agacatagta 50

<210> 5123
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (5124 is other entry)

<221> misc_feature
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<223> Accession number cg43983790

<400> 5123
caactggaaa tactaaacaa atactggaat tcacattaca gacagacgaa a 51

<210> 5124
<211> 51
<212> DNA
<213> Homo sapiens

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<222> (26)...(0)
<223> 2 of 2 allelic variants (5123 is other entry)

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<223> Accession number cg43983790

<400> 5124
caactggaaa tactaaacaa atacttgaat tcacattaca gacagacgaa a 51

<210> 5125
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (5126 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43983790

<400> 5125

cgaaaccaac atggatgcca cacataactt cctttgtagt ttcacagaga g

51

<210> 5126

<211> 51

<212> DNA

<213> Homo sapiens

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<222> (26)...(0)

<223> 2 of 2 allelic variants (5125 is other entry)

<221> misc_feature

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<223> Accession number cg43983790

<400> 5126

cgaaaccaac atggatgcca cacatcactt cctttgtagt ttcacagaga g

51

<210> 5127

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 1 of 2 allelic variants (5128 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43983790

<400> 5127

agagcctatt tgtggttgct caggtggggt catacattgc ttgcagaaat g

51

<210> 5128

<211> 50

<212> DNA

<213> Homo sapiens

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<222> (26)...(0)

<223> 2 of 2 allelic variants (5127 is other entry)

<221> misc_feature

<222> (25)...(26)

<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
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<223> Accession number cg43983790

<400> 5128
agagcctatt tgtggttgct caggtgggtc atacattgct tgcagaaatg 50

<210> 5129
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (5130 is other entry)

<221> misc_feature
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<223> Accession number cg43983790

<400> 5129
gctcaggtgg ggtcatacat tgcttgaga aatggcctga tcatagctct a 51

<210> 5130
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 2 of 2 allelic variants (5129 is other entry)

<221> misc_feature
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<223> Accession number cg43983790

<400> 5130
gctcaggtgg ggtcatacat tgcttacaga aatggcctga tcatagctct a 51

<210> 5131
<211> 51
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<223> 1 of 2 allelic variants (5132 is other entry)

<221> misc_feature
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<223> Accession number cg43983790

<400> 5131
aagttgagat aataatattt cacatattta tatacagaga atcactctca a 51

<210> 5132
<211> 51
<212> DNA
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<223> 2 of 2 allelic variants (5131 is other entry)

<221> misc_feature
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<223> Accession number cg43983790

<400> 5132
aagttgagat aataatattt cacatgttta tatacagaga atcactctca a 51

<210> 5133
<211> 51
<212> DNA
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<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5134 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43983790

<400> 5133
aataggattt gggggtgact tgtacacatt tctaaaaaca cttttctttt t 51

<210> 5134
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<212> DNA
<213> Homo sapiens

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<223> 2 of 2 allelic variants (5133 is other entry)

<221> misc_feature
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<223> Accession number cg43983790

<400> 5134
aataggattt gggggtgact tgtacgcatt tctaaaaaca cttttctttt t 51

<210> 5135
<211> 51
<212> DNA
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<223> 1 of 2 allelic variants (5136 is other entry)

<221> misc_feature
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<223> Accession number cg43983873

<400> 5135
ctactgtcag ttttctgatt gagaacattc atcgcccaag aaacagctca t 51

<210> 5136
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5135 is other entry)

<221> misc_feature
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<223> Accession number cg43983873

<400> 5136
ctactgtcag ttttctgatt gagaatattc atcgcccaag aaacagctca t 51

<210> 5137
<211> 51
<212> DNA
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<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5138 is other entry)

<221> misc_feature
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<223> Accession number cg43983994

<400> 5137
tggcgtggg taccgactca aaggcagcaa tgccattccc tagctcagac a 51

<210> 5138
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (5137 is other entry)

<221> misc_feature

<222> (0)...(0)
<223> Accession number cg43983994

<400> 5138
tggcgggtggg taccgactca aaggcggcaa tgccattccc tagctcagac a 51

<210> 5139
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5140 is other entry)

<221> misc_feature
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<223> Accession number cg43984006

<400> 5139
caccatgcag acacgcagct gtgaacgaca gttcagaact cagcgtaagc t 51

<210> 5140
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5139 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43984006

<400> 5140
caccatgcag acacgcagct gtgaatgaca gttcagaact cagcgtaagc t 51

<210> 5141
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (5142 is other entry)

<221> misc_feature
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<223> Accession number cg43984006

<400> 5141
acgcagctgt gaacgcagct tcagaactca gcgtaagctt gtgctatgaa c 51

<210> 5142
<211> 51
<212> DNA
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<220>
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<223> 2 of 2 allelic variants (5141 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43984006

<400> 5142
acgcagctgt gaacgacagt tcagagctca gcgtaagctt gtgctatgaa c 51

<210> 5143
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<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (5144 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43984006

<400> 5143
gctgtgaacg acagttcaga actcagcgta agcttgtgct atgaacgagc a 51

<210> 5144
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (5143 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43984006

<400> 5144
gctgtgaacg acagttcaga actcaacgta agcttgtgct atgaacgagc a 51

<210> 5145
<211> 51
<212> DNA
<213> Homo sapiens

<220>

<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5146 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43984006

<400> 5145
tgtgaacgac agttcagaac tcagcgtaag cttgtgctat gaacgagcac c 51

<210> 5146
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5145 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43984006

<400> 5146
tgtgaacgac agttcagaac tcagcataag cttgtgctat gaacgagcac c 51

<210> 5147
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5148 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43984006

<400> 5147
aacgacagtt cagaactcag cgtaagcttg tgctatgaac gagcacccgtc a 51

<210> 5148
<211> 51
<212> DNA
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<223> 2 of 2 allelic variants (5147 is other entry)

<221> misc_feature
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<223> Accession number cg43984006

<400> 5148

aacgacagtt cagaactcag cgtaaacttg tgctatgaac gagcaccgtc a

51

<210> 5149

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (5150 is other entry)

<221> misc_feature

<222> (0)...(0)

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<400> 5149

tcagaactca gcgtaagctt gtgctatgaa cgagcaccgt cagagaattc c

51

<210> 5150

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 2 of 2 allelic variants (5149 is other entry)

<221> misc_feature

<222> (0)...(0)

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<400> 5150

tcagaactca gcgtaagctt gtgctgtgaa cgagcaccgt cagagaattc c

51

<210> 5151

<211> 51

<212> DNA

<213> Homo sapiens

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<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (5152 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43984006

<400> 5151

ctcagcgtaa gcttgtgcta tgaacgagca ccgtcagaga attcccaccc a

51

<210> 5152

<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (5151 is other entry)

<221> misc_feature
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<223> Accession number cg43984006

<400> 5152
ctcagcgtaa gcttgtgcta tgaacaagca ccgtcagaga attcccaccc a 51

<210> 5153
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5154 is other entry)

<221> misc_feature
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<223> Accession number cg43984065

<400> 5153
gactccggcg gagcagaagc cttcgttggg ggcggcacag gggtccttaa a 51

<210> 5154
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (5153 is other entry)

<221> misc_feature
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<223> Accession number cg43984065

<400> 5154
gactccggcg gagcagaagc cttcgttggg ggcggcacag gggtccttaa a 51

<210> 5155
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature

<222> (26)...(0)
<223> 1 of 2 allelic variants (5156 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43984242

<400> 5155
gcccacgtcc cagctggacc atggcgcctc gcggaacgt ggtgaagatt g 51

<210> 5156
<211> 50
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (5155 is other entry)

<221> misc_feature
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<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43984242

<400> 5156
gcccacgtcc cagctggacc atggccctcc gcggaacgtg gtgaagattg 50

<210> 5157
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5158 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43984242

<400> 5157
gcggaacgtg gtgaagattg ccatcaagat gcgtagcgcc atcccgcagc t 51

<210> 5158
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5157 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43984242

<400> 5158
gcggaacgtg gtgaagattg ccatccagat gcgtgacgcc atcccgagc t 51

<210> 5159
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5160 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43984508

<400> 5159
cctcgtcgct gtccagcgag gccatctccg tggggtcctc agtggtggcg a 51

<210> 5160
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5159 is other entry)

<221> misc_feature
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<223> Accession number cg43984508

<400> 5160
cctcgtcgct gtccagcgag gccatatccg tggggtcctc agtggtggcg a 51

<210> 5161
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5162 is other entry)

<221> misc_feature
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<223> Accession number cg43984651

<400> 5161

tcctgtttga acttggtgcc aaatagagta actcggactc cagttggagg g 51

<210> 5162

<211> 51

<212> DNA

<213> Homo sapiens

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<222> (26)...(0)

<223> 2 of 2 allelic variants (5161 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43984651

<400> 5162

tcctgtttga acttggtgcc aaataaagta actcggactc cagttggagg g 51

<210> 5163

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (5164 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43984671

<400> 5163

ataccgtgta cactgatata cacgaagctg ctcttcattt ttttgtcaga t 51

<210> 5164

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (5163 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43984671

<400> 5164

ataccgtgta cactgatata cacgaggctg ctcttcattt ttttgtcaga t 51

<210> 5165

<211> 51

<212> DNA

<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5166 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43984751

<400> 5165
aaacactgcc atggtcagcg ggggtggccg aaggggtaac gccccagcag t 51

<210> 5166
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<221> misc_feature
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<400> 5166
aaacactgcc atggtcagcg ggggttgccg aaggggtaac gccccagcag t 51

<210> 5167
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<400> 5167
cactgccatg gtcagcgggg gtggccgaag gggtaacgcc ccagcagtcc g 51

<210> 5168
<211> 51
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<400> 5168
cactgccatg gtcagcgggg gtggctgaag gggtaacgcc ccagcagtc g 51

<210> 5169
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ggtcagcggg ggtggccgaa ggggtaacgc cccagcagtc cgaggccagc t 51

<210> 5170
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<210> 5171
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<223> 1 of 2 allelic variants (5172 is other entry)

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aggggtaacg cccagcagtc ccgaggccag ctccgagggt ctaacttctc c 51

<210> 5172
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<223> 2 of 2 allelic variants (5171 is other entry)

<221> misc_feature
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aggggtaacg cccagcagc cgcagaccag ctccgagggt ctaacttctc c 51

<210> 5173
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<223> 1 of 2 allelic variants (5174 is other entry)

<221> misc_feature
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<223> Accession number cg43984865

<400> 5173
acacatcagg gtaaatgggg tatccatgac ctcaagcacc cctcacaaaa t 51

<210> 5174
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 2 of 2 allelic variants (5173 is other entry)

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<400> 5174
acacatcagg gtaaatgggg tatccctgac ctcaagcacc cctcacaaaa t 51

<210> 5175
<211> 51
<212> DNA
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<221> misc_feature
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ccatgacctc aagcaccct cacaaaatct gagtactcac aagatctggt t 51

<210> 5176
<211> 51
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<223> 2 of 2 allelic variants (5175 is other entry)

<221> misc_feature
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<400> 5176
ccatgacctc aagcaccct cacaagatct gagtactcac aagatctggt t 51

<210> 5177
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<212> DNA
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<223> 1 of 2 allelic variants (5178 is other entry)

<221> misc_feature
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<400> 5177
atgacctcaa gcaccctca caaaatctga gtactcaca gatctggttg t 51

<210> 5178
<211> 51
<212> DNA
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<223> 2 of 2 allelic variants (5177 is other entry)

<221> misc_feature

<222> (0)...(0)
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<400> 5178
atgacctcaa gcacctca caaacctga gtactcacia gatctggtg t 51

<210> 5179
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (5180 is other entry)

<221> misc_feature
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<223> Accession number cg43984962

<400> 5179
acgtctgtga aagctaata tccacagtga tgttccggaa aaggaagcc a 51

<210> 5180
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<212> DNA
<213> Homo sapiens

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<223> 2 of 2 allelic variants (5179 is other entry)

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<223> Accession number cg43984962

<400> 5180
acgtctgtga aagctaata tccactgtga tgttccggaa aaggaagcc a 51

<210> 5181
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 1 of 2 allelic variants (5182 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43985092

<400> 5181
caggagggaa.gaggccgctg ggccccagga atcctaggcc tctcttcccg g 51

<210> 5182
<211> 50
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<221> misc_feature
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<400> 5182
caggaggggaa gaggccgctg ggcccaggca tcctaggcct ctcttcccg

50

<210> 5183
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (5184 is other entry)

<221> misc_feature
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<223> Accession number cg43985377

<400> 5183
ataataattt cagggaaaaa aaaaaaaggc acatccaggc accacattca a

51

<210> 5184
<211> 50
<212> DNA
<213> Homo sapiens

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<223> 2 of 2 allelic variants (5183 is other entry)

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<223> Accession number cg43985377

<400> 5184

ataataatattt cagggaaaaa aaaaaaggca catccaggca ccacattcaa

50

<210> 5185

<211> 51

<212> DNA

<213> Homo sapiens

<220>

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<222> (26)...(0)

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<223> Accession number cg43985471

<400> 5185

gccctccag gcagctgggg gagagcggag gcagggtcgc tcagggttct a

51

<210> 5186

<211> 51

<212> DNA

<213> Homo sapiens

<220>

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<222> (26)...(0)

<223> 2 of 2 allelic variants (5185 is other entry)

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<223> Accession number cg43985471

<400> 5186

gccctccag gcagctgggg gagagtggag gcagggtcgc tcagggttct a

51

<210> 5187

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 1 of 2 allelic variants (5188 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43985734

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ggtgaagagt gtctgatgcc agcgccatc ttggaccttg cagccatcca t

51

<210> 5188

<211> 50

<212> DNA

<213> Homo sapiens

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<223> 2 of 2 allelic variants (5187 is other entry)

<221> misc_feature
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<221> misc_feature
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<223> Accession number cg43985734

<400> 5188
ggtgaagagt gtctgatgcc agcggcatct tggaccttgc agccatccat 50

<210> 5189
<211> 51
<212> DNA
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<223> 1 of 2 allelic variants (5190 is other entry)

<221> misc_feature
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<223> Accession number cg43985772

<400> 5189
atggactcag cctgggtcag agagggaact tctgaagcta cacgaacaag c 51

<210> 5190
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (5189 is other entry)

<221> misc_feature
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<223> Accession number cg43985772

<400> 5190
atggactcag cctgggtcag agaggaaact tctgaagcta cacgaacaag c 51

<210> 5191
<211> 51
<212> DNA
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<223> 1 of 2 allelic variants (5192 is other entry)

<221> misc_feature
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<223> Accession number cg43986072

<400> 5191
tatcagtaaa cattaatttt ttttttcctt gaggcacagc atgatcttgg c 51

<210> 5192
<211> 50
<212> DNA
<213> Homo sapiens

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<400> 5192
tatcagtaaa cattaatttt ttttttccttg aggcacagca tgatcttggc 50

<210> 5193
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<223> 1 of 2 allelic variants (5194 is other entry)

<221> misc_feature
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<223> Accession number cg43986256

<400> 5193
cttctgagga tcccaggaca gggccaccag caggggaggc cctcaatacc t 51

<210> 5194
<211> 51
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<220>
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<223> 2 of 2 allelic variants (5193 is other entry)

<221> misc_feature

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<223> Accession number cg43986256

<400> 5194

cttctgagga tcccaggaca gggccgccag caggggaggc cctcaatacc t

51

<210> 5195

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 1 of 2 allelic variants (5196 is other entry)

<221> misc_feature

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<223> Accession number cg43986256

<400> 5195

tcaataacctg agcccaaaca acaatgagca gattcccagg ctttgctttc a

51

<210> 5196

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 2 of 2 allelic variants (5195 is other entry)

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<223> Accession number cg43986256

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51

<210> 5197

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 1 of 2 allelic variants (5198 is other entry)

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<222> (0)...(0)

<223> Accession number cg43986313

<400> 5197
cagagatcct gcaggagttg gagtatggcc cctccgtgga ctggtgggcc c 51

<210> 5198
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<223> 2 of 2 allelic variants (5197 is other entry)

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<400> 5198
cagagatcct gcaggagttg gactacggcc cctccgtgga ctggtgggcc c 51

<210> 5199
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<223> 1 of 2 allelic variants (5200 is other entry)

<221> misc_feature
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<400> 5199
ctgtgtggca tcgcagaatg gcgaggacgc catcaagcag cacccattct t 51

<210> 5200
<211> 51
<212> DNA
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<400> 5200
ctgtgtggca tcgcagaatg gcgagaacgc catcaagcag cacccattct t 51

<210> 5201
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<213> Homo sapiens

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<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (5202 is other entry)

<221> misc_feature

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<400> 5201

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<210> 5202

<211> 51

<212> DNA

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<222> (0)...(0)

<223> Accession number cg43986510

<400> 5202

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<210> 5203

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<212> DNA

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<223> 1 of 2 allelic variants (5204 is other entry)

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<223> Accession number cg43986519

<400> 5203

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<210> 5204

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<213> Homo sapiens

<220>

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<222> (26)...(0)

<223> 2 of 2 allelic variants (5203 is other entry)

<221> misc_feature
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<223> Accession number cg43986519

<400> 5204
taattttaaca gtgacagaac gacagacagt gtccataagc aaacaacaca c 51

<210> 5205
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 1 of 2 allelic variants (5206 is other entry)

<221> misc_feature
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<223> Accession number cg43986519

<400> 5205
gacagcagaa ttaagattta aaaaacgaaa gcatttcac c aaaacaaaaa a 51

<210> 5206
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (5205 is other entry)

<221> misc_feature
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<223> Accession number cg43986519

<400> 5206
gacagcagaa ttaagattta aaaaatgaaa gcatttcac c aaaacaaaaa a 51

<210> 5207
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<212> DNA
<213> Homo sapiens

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<223> 1 of 2 allelic variants (5208 is other entry)

<221> misc_feature
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<222> (0)...(0)
<223> Accession number cg43986519

<400> 5207
gaaagcattt catcaaaaca aaaaactaac ttcttcacag gaagtccact 50

<210> 5208
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 2 of 2 allelic variants (5207 is other entry)

<221> misc_feature
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<223> Accession number cg43986519

<400> 5208
gaaagcattt catcaaaaca aaaaaactaa ctcttcaca ggaagtccac t 51

<210> 5209
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<223> 1 of 2 allelic variants (5210 is other entry)

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<223> Accession number cg43986639

<400> 5209
tggcgccac ttttctcca caatctgaac acgcccttag cttaactgca g 51

<210> 5210
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (5209 is other entry)

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<400> 5210
tggcgccac ttttctcca caatccgaac acgcccttag cttaactgca g 51

<210> 5211
<211> 51
<212> DNA
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<223> 1 of 2 allelic variants (5212 is other entry)

<221> misc_feature
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<223> Accession number cg43986776

<400> 5211
agagctccag ggttcagaa ggcgtgcac aggtcacgaa ggaatggctg t 51

<210> 5212
<211> 50
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
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<221> misc_feature
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<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
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<400> 5212
agagctccag ggttcagaa ggcgtgcaca gggtcacgaag gaatggctgt 50

<210> 5213
<211> 51
<212> DNA
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5214 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43986966

<400> 5213
tagcaatatt gttaatcact gctcttattt tggagacaaa aggaccaact t 51

<210> 5214
<211> 51

<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5213 is other entry)

<221> misc_feature
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<400> 5214
tagcaatatc gttaatcact gctctgattt tggagacaaa aggaccaact t 51

<210> 5215
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5216 is other entry)

<221> misc_feature
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<223> Accession number cg43986974

<400> 5215
atggtgaatt gggaatgaag ctgtcggaaa tccccttgac tctgcattct g 51

<210> 5216
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<400> 5216
atggtgaatt gggaatgaag ctgtcagaaa tccccttgac tctgcattct g 51

<210> 5217
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<220>
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<223> 1 of 2 allelic variants (5218 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43987185

<400> 5217

cacaaatgtc agaacattaa agtagcattt tttccatagg aatatataaa a

51

<210> 5218

<211> 51

<212> DNA

<213> Homo sapiens

<220>

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<222> (26)...(0)

<223> 2 of 2 allelic variants (5217 is other entry)

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<222> (0)...(0)

<223> Accession number cg43987185

<400> 5218

cacaaatgtc agaacattaa agtagtattt tttccatagg aatatataaa a

51

<210> 5219

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<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (5220 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43987473

<400> 5219

ccgttacaac aagcttcctt gatgggtgcc acagctttta caaatgctgg t

51

<210> 5220

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 2 of 2 allelic variants (5219 is other entry)

<221> misc_feature

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<223> Accession number cg43987473

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<210> 5221
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<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5222 is other entry)

<221> misc_feature
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<223> Accession number cg43987479

<400> 5221
atcagacaga catgacctcc tacgcagcac cagtatgcat gagcgggtga a 51

<210> 5222
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<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (5221 is other entry)

<221> misc_feature
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<223> Accession number cg43987479

<400> 5222
atcagacaga catgacctcc tacgcggcac cagtatgcat gagcgggtga a 51

<210> 5223
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5224 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43987639

<400> 5223
ataataaatc gtttgcgttt cctaagaaac tgggtttttc ttttccttg t 51

<210> 5224
<211> 51
<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (5223 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43987639

<400> 5224

ataataaatc gtttgcgttt cctaaaaaac tgggtttttc ttttccttg t

51

<210> 5225

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (5226 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43987678

<400> 5225

cccgagctg aagatggggc agaagcccag gcccctctc cacccccagc t

51

<210> 5226

<211> 50

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

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<223> 2 of 2 allelic variants (5225 is other entry)

<221> misc_feature

<222> (25)...(26)

<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43987678

<400> 5226

cccgagctg aagatggggc agaagccagg ccccatctcc acccccagct

50

<210> 5227

<211> 51

<212> DNA

<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (5228 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43987682

<400> 5227
agaactgcag ggtgttctca ctaaacagtg gccctgccag ctggcggatc a 51

<210> 5228
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (5227 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43987682

<400> 5228
agaactgcag ggtgttctca ctaaatagtg gccctgccag ctggcggatc a 51

<210> 5229
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5230 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43987769

<400> 5229
taaagttaat taccttacct ttgcaactat tttctgtata agaattctca a 51

<210> 5230
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (5229 is other entry)

<221> misc_feature

<222> (0)...(0)
<223> Accession number cg43987769

<400> 5230
taaagttaat taccttacct ttgcagctat tttctgtata agaattctcaa a 51

<210> 5231
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5232 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43987787

<400> 5231
actgaaatat atagaaaacc ccaatgtatg aaacaagttt taggcattgg t 51

<210> 5232
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5231 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43987787

<400> 5232
actgaaatat atagaaaacc ccaatatatg aaacaagttt taggcattgg t 51

<210> 5233
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5234 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43987853

<400> 5233
aatcttagtt ttactatata ttatttggtg ctcttggtt cagtagttgg g 51

<210> 5234
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5233 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43987853

<400> 5234
aatcttagtt ttactatata ttattcggta ctctggatt cagtagttgg g 51

<210> 5235
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5236 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43987853

<400> 5235
ggattcagta gttggggtaa aaaaagtttc tatattcaaa catatctaca g 51

<210> 5236
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (5235 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43987853

<400> 5236
ggattcagta gttggggtaa aaaaaatttc tatattcaaa catatctaca g 51

<210> 5237
<211> 51
<212> DNA
<213> Homo sapiens

<220>

<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5238 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43987853

<400> 5237
agtagttggg gtaaaaaaag tttctatatt caaacatatc tacagaatgt c 51

<210> 5238
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5237 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43987853

<400> 5238
agtagttggg gtaaaaaaag tttctgtatt caaacatatc tacagaatgt c 51

<210> 5239
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5240 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43987905

<400> 5239
ggcagaaagc gcaccagctt cagcacacgc agcagccgga aggtgcgcag c 51

<210> 5240
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5239 is other entry)

<221> misc_feature
<222> (0)...(0)

<223> Accession number cg43987905

<400> 5240

ggcagaaagc gcaccagctt cagcatacgc agcagccgga aggtgcgcag c

51

<210> 5241

<211> 51

<212> DNA

<213> Homo sapiens

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<222> (26)...(0)

<223> 1 of 2 allelic variants (5242 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43988015

<400> 5241

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51

<210> 5242

<211> 51

<212> DNA

<213> Homo sapiens

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<222> (26)...(0)

<223> 2 of 2 allelic variants (5241 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43988015

<400> 5242

accttggcct cccaaagtgc tgggactaca ggcatgagca accgcaccgc g

51

<210> 5243

<211> 51

<212> DNA

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<223> 1 of 2 allelic variants (5244 is other entry)

<221> misc_feature

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<223> Accession number cg43988019

<400> 5243

gtggtagaga catgattggt attgccaaaa caggtagtgg gaaaactgca g

51

<210> 5244

<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5243 is other entry)

<221> misc_feature
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<223> Accession number cg43988019

<400> 5244
gtggtagaga catgattggt attgcaaaaa caggtagtgg gaaaactgca g 51

<210> 5245
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5246 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43988355

<400> 5245
ggttttgctg ggacagcctg tgaactctgt gctcctggtg cctttgggcc c 51

<210> 5246
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5245 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43988355

<400> 5246
ggttttgctg ggacagcctg tgaacgctgt gctcctggtg cctttgggcc c 51

<210> 5247
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature

<222> (26)...(0)
<223> 1 of 2 allelic variants (5248 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43988355

<400> 5247
actgtagcca ggtaggaaca atgggtcactt gtacctgcct gcccgactac g 51

<210> 5248
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (5247 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43988355

<400> 5248
actgtagcca ggtaggaaca atgggttactt gtacctgcct gcccgactac g 51

<210> 5249
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5250 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43988355

<400> 5249
atcttgccct tttccaagcg ggtgatccgg tcccactgaa gtgagccctt c 51

<210> 5250
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5249 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43988355

<400> 5250
atcttgccct tttccaagcg ggtgaccg tcccaactgaa gtgagccctt c 51

<210> 5251
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5252 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43988361

<400> 5251
ccatgacggt gcgctcgggg ttgatgccgt attcctggga cacgcagtcg a 51

<210> 5252
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (5251 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43988361

<400> 5252
ccatgacggt gcgctcgggg ttgataccgt attcctggga cacgcagtcg a 51

<210> 5253
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5254 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43988361

<400> 5253
tgacggtgcg ctcgggggtg atgccgtatt cctgggacac gcagtcgaaa a 51

<210> 5254
<211> 51

<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (5253 is other entry)

<221> misc_feature
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<223> Accession number cg43988361

<400> 5254
tgacggtgcg ctcggggttg atgccatatt cctgggacac gcagtcgaaa a 51

<210> 5255
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5256 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43988361

<400> 5255
agcggctggg cttcccgatg atgtccgcct ggcgctgggc ggccatctcc a 51

<210> 5256
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (5255 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43988361

<400> 5256
agcggctggg cttcccgatg atgtccgcct ggcgctgggc ggccatctcc a 51

<210> 5257
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)

<223> 1 of 2 allelic variants (5258 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43988588

<400> 5257

caaaaacagc taaagcagca aattccaagt caaaagagag tgatgaacct c

51

<210> 5258

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (5257 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43988588

<400> 5258

caaaaacagc taaagcagca aattcaaagt caaaagagag tgatgaacct c

51

<210> 5259

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (5260 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43988588

<400> 5259

aaagcgcagg cccagctcca gaagcgctac aaagaccttg aacaacaaga c

51

<210> 5260

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (5259 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43988588

<400> 5260
aaagcgcagg cccagctcca gaagcactac aaagaccttg aacaacaaga c 51

<210> 5261
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5262 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43988624

<400> 5261
cccagagcct gcgtttcttg gcacagacac agagagaaat ggaataaatt a 51

<210> 5262
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (5261 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43988624

<400> 5262
cccagagcct gcgtttcttg gcacacacac agagagaaat ggaataaatt a 51

<210> 5263
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5264 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43988652

<400> 5263
gctccacttt gcaggtgacc gtgacacat ctgctcggcc caagggcacc a 51

<210> 5264
<211> 51
<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (5263 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43988652

<400> 5264

gtccacttt gcaggtgacc gtgacgcat ctgctcgcc caagggcacc a

51

<210> 5265

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (5266 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43988652

<400> 5265

ctttgcaggt gaccgtgaca ccatctgctc ggccaaggg caccaaggtg a

51

<210> 5266

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (5265 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43988652

<400> 5266

ctttgcaggt gaccgtgaca ccatccgctc ggccaaggg caccaaggtg a

51

<210> 5267

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (5268 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43988747

<400> 5267
agaggaatcg gtccacagcc ttccgggttg tgggtgtaag tgccagcact t 51

<210> 5268
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (5267 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43988747

<400> 5268
agaggaatcg gtccacagcc ttccgagttg tgggtgtaag tgccagcact t 51

<210> 5269
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5270 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43989086

<400> 5269
tgccactcca ggtgggggga gtggtgcccc agccacgctt caacccttct c 51

<210> 5270
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (5269 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43989086

<400> 5270

tgccactcca ggtgggggga gtggtacccc agccacgctt caacccttct c 51

<210> 5271
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5272 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43989459

<400> 5271
gcaggtgtat ctgcacagtg gtcgccccac agcagaccat gtgttcacgg g 51

<210> 5272
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (5271 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43989459

<400> 5272
gcaggtgtat ctgcacagtg gtcgctccac agcagaccat gtgttcacgg g 51

<210> 5273
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5274 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43990145

<400> 5273
ccttgaattc cagaagaagg aagaaacctt atgtctagca gtctgagcct g 51

<210> 5274
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (5273 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43990145

<400> 5274
ccttgaattc cagaagaagg aagaagcctt atgtctagca gtctgagcct g 51

<210> 5275
<211> 50
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5276 is other entry)

<221> misc_feature
<222> (25)...(26)
<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43990145

<400> 5275
ccctgtctct agaaagaaaa aaaaaggaaac tgtgtgagag gaaattcatg 50

<210> 5276
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (5275 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43990145

<400> 5276
ccctgtctct agaaagaaaa aaaaaaggaa ctgtgtgaga ggaaattcat g 51

<210> 5277
<211> 51
<212> DNA
<213> Homo sapiens

<220>

<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5278 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43990709

<400> 5277
cacgttttcc actgacataa agttgcttcg ccccttgcag cttatctcca c 51

<210> 5278
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (5277 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43990709

<400> 5278
cacgttttcc actgacataa agttgtttcg ccccttgcag cttatctcca c 51

<210> 5279
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5280 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43990709

<400> 5279
tttccactga cataaagttg cttcgccct tgcagcttat ctccaccttc a 51

<210> 5280
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> Accession number cg43990709

<400> 5280

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51

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<223> 1 of 2 allelic variants (5282 is other entry)

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<223> Accession number cg43991361

<400> 5281

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51

<210> 5282

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<223> Accession number cg43991361

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51

<210> 5283

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<212> DNA

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<223> 1 of 2 allelic variants (5284 is other entry)

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<223> Accession number cg43991427

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51

<210> 5284

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<210> 5285
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<223> 1 of 2 allelic variants (5286 is other entry)

<221> misc_feature
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<223> Accession number cg43991837

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gagtatcaac tacacatttt ttttttcaag cagcatatta ttataaagcc c 51

<210> 5286
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gagtatcaac tacacatttt tttttcaagc agcatattat tataaagccc 50

<210> 5287
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<220>

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<223> 1 of 2 allelic variants (5288 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43991837

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ttttaattgc ttttttagga aaaagaaaaa aaaagggtgc ttttaatact t

51

<210> 5288

<211> 50

<212> DNA

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50

<210> 5289

<211> 51

<212> DNA

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<223> Accession number cg43992019

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cggaagcttt tttctcagga ggtaatctct aataacaagc agagtgcct c

51

<210> 5290

<211> 51

<212> DNA

<213> Homo sapiens

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<223> Accession number cg43992019

<400> 5290
cggaagcttt tttctcagga ggtaaactct aataacaagc agagtgcct c 51

<210> 5291
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<212> DNA
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<223> Accession number cg43992019

<400> 5291
atcccaacac acagccagtc aacgagcctc tggccccttc ctctgggtcc t 51

<210> 5292
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<400> 5292
atcccaacac acagccagtc aacgacctct ggccccttcc tctgggtcct 50

<210> 5293
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<223> Accession number cg43992029

<400> 5293
ggggcagcta cagggttcag ctctgggcag ggcttggcca gggacagtgt g 51

<210> 5294
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 2 of 2 allelic variants (5293 is other entry)

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<223> Accession number cg43992029

<400> 5294
ggggcagcta cagggttcag ctctgagcag ggcttggcca gggacagtgt g 51

<210> 5295
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<223> Accession number cg43992029

<400> 5295
cgcacagcca gacacacaca cacacaccct gccacgcaca gcacgcaggc a 51

<210> 5296
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<223> 2 of 2 allelic variants (5295 is other entry)

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<223> Accession number cg43992029

<400> 5296
cgcacagcca gacacacaca cacacccctg ccacgcacag cacgcaggca 50

<210> 5297
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<223> 1 of 2 allelic variants (5298 is other entry)

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<400> 5297
cgcacagcac gcaggcacac acacacttgt gcatgcacac gcgttcatat a 51

<210> 5298
<211> 50
<212> DNA
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<223> 2 of 2 allelic variants (5297 is other entry)

<221> misc_feature
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<221> misc_feature
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<223> Accession number cg43992029

<400> 5298
cgcacagcac gcaggcacac acacattgtg catgcacacg cgttcatata 50

<210> 5299
<211> 51
<212> DNA
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<223> 1 of 2 allelic variants (5300 is other entry)

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<222> (0)...(0)
<223> Accession number cg43992178

<400> 5299
ataaacacgt aagagtaaca ctttgcactc caatagcacc tgttggtcaa a 51

<210> 5300
<211> 51
<212> DNA
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<223> 2 of 2 allelic variants (5299 is other entry)

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<223> Accession number cg43992178

<400> 5300
ataaacacgt aagagtaaca ctttgtactc caatagcacc tgttggtcaa a 51

<210> 5301
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<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (5302 is other entry)

<221> misc_feature
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<223> Accession number cg43992186

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tggccttggt aaaggaaatg acacacattc ccccaatttg gaagcaatct t 51

<210> 5302
<211> 51
<212> DNA
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<223> 2 of 2 allelic variants (5301 is other entry)

<221> misc_feature
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<223> Accession number cg43992186

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<223> 1 of 2 allelic variants (5304 is other entry)

<221> misc_feature
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<223> Accession number cg43992349

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caggagagct gcctcattag acttcgccag tggctccagg agaccacaa g 51

<210> 5304
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<223> 2 of 2 allelic variants (5303 is other entry)

<221> misc_feature
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<400> 5304
caggagagct gcctcattag acttcaccag tggctccagg agaccacaa g 51

<210> 5305
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<212> DNA
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<223> 1 of 2 allelic variants (5306 is other entry)

<221> misc_feature
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<223> Accession number cg43992349

<400> 5305
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<210> 5306
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<223> 2 of 2 allelic variants (5305 is other entry)

<221> misc_feature
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<223> Accession number cg43992349

<400> 5306
gagagctgcc tcattagact tcgccggtgg ctccaggaga cccacaaggg c 51

<210> 5307
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<223> 1 of 2 allelic variants (5308 is other entry)

<221> misc_feature
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<223> Accession number cg43992349

<400> 5307
cagagagatc atgtgtcagt ctttgacgtg gagaaagcag catcaggtag a 51

<210> 5308
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<223> 2 of 2 allelic variants (5307 is other entry)

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<223> Accession number cg43992349

<400> 5308
cagagagatc atgtgtcagt ctttggcgtg gagaaagcag catcaggtag a 51

<210> 5309
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5310 is other entry)

<221> misc_feature
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<223> Accession number cg43992895

<400> 5309

gcaccaccac acccgctaa ttttttatat ctttagcaga gatggggttt c

51

<210> 5310

<211> 51

<212> DNA

<213> Homo sapiens

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<222> (26)...(0)

<223> 2 of 2 allelic variants (5309 is other entry)

<221> misc_feature

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<223> Accession number cg43992895

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51

<210> 5311

<211> 51

<212> DNA

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<222> (26)...(0)

<223> 1 of 2 allelic variants (5312 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43993206

<400> 5311

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51

<210> 5312

<211> 51

<212> DNA

<213> Homo sapiens

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<223> Accession number cg43993206

<400> 5312

atgtgtagag tagattgtct ggtgccctca gttgttttta tttacatttg t

51

<210> 5313

<211> 51
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<223> 1 of 2 allelic variants (5314 is other entry)

<221> misc_feature
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<223> Accession number cg43993206

<400> 5313
ctaaccacgc gtacgacaac tggtagaaaac aaactcaagt gcaaaggaaa c 51

<210> 5314
<211> 51
<212> DNA
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<223> 2 of 2 allelic variants (5313 is other entry)

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ctaaccacgc gtacgacaac tggtagaaaac aaactcaagt gcaaaggaaa c 51

<210> 5315
<211> 51
<212> DNA
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<223> Accession number cg43993218

<400> 5315
cttttctgtc agcaaaaaag gtacaatttt tttaaaactt gaaaatcaat a 51

<210> 5316
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<400> 5316
cttttctgtc agcaaaaaag gtacattttt tttaaaactt gaaaatcaat a 51

<210> 5317
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<210> 5318
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<212> DNA
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<221> misc_feature
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tgaagcagcc cgaaaacaag tgaaaaggaa accaagataa ctctctctcc c 51

<210> 5319
<211> 51
<212> DNA
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<223> 1 of 2 allelic variants (5320 is other entry)

<221> misc_feature
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<223> Accession number cg43993836

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<210> 5320
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<212> DNA
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<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5319 is other entry)

<221> misc_feature
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<223> Accession number cg43993836

<400> 5320
ataggccggg caagcggcac ccaggagagc ctcgacacag gtagtgacct g 51

<210> 5321
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5322 is other entry)

<221> misc_feature
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<223> Accession number cg43993844

<400> 5321
gtgcacttct cgccgggggc ccgggcgctc accacacctg cagtgcattg c 51

<210> 5322
<211> 51
<212> DNA
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<223> 2 of 2 allelic variants (5321 is other entry)

<221> misc_feature
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<223> Accession number cg43993844

<400> 5322

gtgcacttct cgccgggggc ccgggggctc accacacctg cagtgcattgt c 51

<210> 5323

<211> 51

<212> DNA

<213> Homo sapiens

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<222> (26)...(0)

<223> 1 of 2 allelic variants (5324 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43993844

<400> 5323

gtggctgatg gtggacactg tcataggaga aggagagttt ggggaagtgt a 51

<210> 5324

<211> 51

<212> DNA

<213> Homo sapiens

<220>

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<222> (26)...(0)

<223> 2 of 2 allelic variants (5323 is other entry)

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<222> (0)...(0)

<223> Accession number cg43993844

<400> 5324

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<210> 5325

<211> 51

<212> DNA

<213> Homo sapiens

<220>

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<223> 1 of 2 allelic variants (5326 is other entry)

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<222> (0)...(0)

<223> Accession number cg43993844

<400> 5325

ggtgcacccc gcatgcactg cagccctgat ggcgagtggc tggcgacctgt a 51

<210> 5326

<211> 50

<212> DNA

<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (5325 is other entry)

<221> misc_feature
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<221> misc_feature
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<223> Accession number cg43993844

<400> 5326
gggtgcacccc gcatgcactg cagcctgatg gcgagtggct ggtgcctgta 50

<210> 5327
<211> 51
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<220>
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<223> 1 of 2 allelic variants (5328 is other entry)

<221> misc_feature
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<223> Accession number cg43993862

<400> 5327
aggctggagt gaagtggcac gatcccgggt cactgcaacc tctgcctccc g 51

<210> 5328
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (5327 is other entry)

<221> misc_feature
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<223> Accession number cg43993862

<400> 5328
aggctggagt gaagtggcac gatcctgggt cactgcaacc tctgcctccc g 51

<210> 5329
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<220>

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<223> 1 of 2 allelic variants (5330 is other entry)

<221> misc_feature
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<223> Accession number cg43993862

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cctcggcctc ccaaagtgc agcattacag gcgtgagcca ccatgcctgg c 51

<210> 5330
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5329 is other entry)

<221> misc_feature
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<223> Accession number cg43993862

<400> 5330
cctcggcctc ccaaagtgc agcatcacag gcgtgagcca ccatgcctgg c 51

<210> 5331
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5332 is other entry)

<221> misc_feature
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<223> Accession number cg43994222

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gagaaacgca ggttgctgcg tgggtaaaaa aaatatttgg agatcatcct a 51

<210> 5332
<211> 50
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5331 is other entry)

<221> misc_feature
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<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43994222

<400> 5332

gagaaacgca ggttgctgcg tggttaaaaa aatatttggga gatcatccta

50

<210> 5333

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

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<223> 1 of 2 allelic variants (5334 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43994222

<400> 5333

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51

<210> 5334

<211> 50

<212> DNA

<213> Homo sapiens

<220>

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<222> (26)...(0)

<223> 2 of 2 allelic variants (5333 is other entry)

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<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43994222

<400> 5334

gcaggttgct gcgtgggttaa aaaaatattt ggagatcatc ctattccaca

50

<210> 5335

<211> 51

<212> DNA

<213> Homo sapiens

<220>

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<222> (26)...(0)

<223> 1 of 2 allelic variants (5336 is other entry)

<221> misc_feature
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<223> Accession number cg43994222

<400> 5335
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<210> 5336
<211> 51
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<220>
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<223> 2 of 2 allelic variants (5335 is other entry)

<221> misc_feature
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<400> 5336
cgagtcagaa gccaaagtatc ttcaaaacct tctcatggag agtgtgaatt t 51

<210> 5337
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (5338 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43994840

<400> 5337
cacagcaagt ccgatgagtc cggccagtcc cgccacaccc agggccatgc c 51

<210> 5338
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (5337 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43994840

<400> 5338
cacagcaagt ccgatgagtc cggcctgtcc cgccacaccc agggccatgc c 51

<210> 5339
<211> 50
<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (5340 is other entry)

<221> misc_feature
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<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43994860

<400> 5339
tttgaagaa ataaaaagaa aaaaatgaag tctcttgtct caatgttgaa 50

<210> 5340
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 2 of 2 allelic variants (5339 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43994860

<400> 5340
tttgaagaa ataaaaagaa aaaaatgaa gtctctgtgc tcaatgttga a 51

<210> 5341
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5342 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43994987

<400> 5341
cttgaaaagg tgctaagatt ggtttctgtt aacatcaaaa aaaaaaaca g 51

<210> 5342

<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (5341 is other entry)

<221> misc_feature
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<223> Accession number cg43994987

<400> 5342
cttgaaaagg tgctaagatt ggtttttggtt aacatcaaaa aaaaaaaaca g 51

<210> 5343
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<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (5344 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43995124

<400> 5343
aagaggcttt gaaaatgtag aactgggagt cataggaaaa aagaagaaag t 51

<210> 5344
<211> 50
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (5343 is other entry)

<221> misc_feature
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<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43995124

<400> 5344
aagaggcttt gaaaatgtag aactgggagtc ataggaaaaa agaagaaagt 50

<210> 5345
<211> 51
<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (5346 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43995124

<400> 5345

agaggctttg aaaatgtaga actgggagtc ataggaaaaa agaagaaagt c

51

<210> 5346

<211> 50

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

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<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43995124

<400> 5346

agaggctttg aaaatgtaga actggagtca taggaaaaaa gaagaaagtc

50

<210> 5347

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (5348 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43995294

<400> 5347

aaaaaggtag atcagaaaca gaaacatgct actatcaggg caactgagct t

51

<210> 5348

<211> 51

<212> DNA

<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (5347 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43995294

<400> 5348
aaaaaggtac atcagaaaca gaaacgtgct actatcaggg caactgagct t 51

<210> 5349
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (5350 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43995376

<400> 5349
gccatgtctg tggtagaggg tgagtaagag gccagagctg agggtagaggt g 51

<210> 5350
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5349 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43995376

<400> 5350
gccatgtctg tggtagaggg tgagtcagag gccagagctg agggtagaggt g 51

<210> 5351
<211> 51
<212> DNA
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<220>
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<223> 1 of 2 allelic variants (5352 is other entry)

<221> misc_feature

<222> (0)...(0)
<223> Accession number cg43995405

<400> 5351
ctgcggtggc cagggccgtg agtcccgtgg cagagccttc tgggcgctgc g 51

<210> 5352
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5351 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43995405

<400> 5352
ctgcggtggc cagggccgtg agtcctgtgg cagagccttc tgggcgctgc g 51

<210> 5353
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5354 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43995405

<400> 5353
ctctcggtcc ggaacaagac gcctcagcca cggctcccc tcggcctatt a 51

<210> 5354
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (5353 is other entry)

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<223> Accession number cg43995405

<400> 5354
ctctcggtcc ggaacaagac gcctcggcca cggctcccc tcggcctatt a 51

<210> 5355
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5356 is other entry)

<221> misc_feature
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<223> Accession number cg43995405

<400> 5355
tacattgccca gctggaaggg cctgggcagg tttctgaaca gcctgggcac c 51

<210> 5356
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (5355 is other entry)

<221> misc_feature
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<400> 5356
tacattgccca gctggaaggg cctggccagg tttctgaaca gcctgggcac c 51

<210> 5357
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (5358 is other entry)

<221> misc_feature
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<223> Accession number cg43995405

<400> 5357
acgcctcgct ggccgcctac caccctggg tcgtgcgcg tgccgtcacc g 51

<210> 5358
<211> 50
<212> DNA
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<220>

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<223> 2 of 2 allelic variants (5357 is other entry)

<221> misc_feature
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<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43995405

<400> 5358
acgcctcgct ggccgcctac caccctgggt cgtgcgccgt gccgtcacg 50

<210> 5359
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5360 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43995405

<400> 5359
cctaccaccc ctgggtcggtg cgccgtgccg tcaccgtggc cttctgcacg c 51

<210> 5360
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5359 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43995405

<400> 5360
cctaccaccc ctgggtcggtg cgccgtgccg tcaccgtggc cttctgcacg c 51

<210> 5361
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)

<223> 1 of 2 allelic variants (5362 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43995762

<400> 5361

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51

<210> 5362

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 2 of 2 allelic variants (5361 is other entry)

<221> misc_feature

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<223> Accession number cg43995762

<400> 5362

gcaccctaac accccgactg gtgttcctgg gtcctaaca ttgtctgtac c

51

<210> 5363

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 1 of 2 allelic variants (5364 is other entry)

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<222> (0)...(0)

<223> Accession number cg43995779

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<210> 5364

<211> 51

<212> DNA

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<223> 2 of 2 allelic variants (5363 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43995779

<400> 5364
aaggactgcc ccacaggaaa cctgaccgtg gacgagttca agaagatcta c 51

<210> 5365
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5366 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43996227

<400> 5365
cccattctata tacagatgtg gcatggctct gccctggcac agccagctgg c 51

<210> 5366
<211> 50
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (5365 is other entry)

<221> misc_feature
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<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43996227

<400> 5366
cccattctata tacagatgtg gcatgctctg ccctggcaca gccagctggc 50

<210> 5367
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5368 is other entry)

<221> misc_feature
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<223> Accession number cg43996227

<400> 5367
tgtgtcagge cagggagtcc ctggcctgcc ctggatagag tggagggccc t 51

<210> 5368
<211> 50
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (5367 is other entry)

<221> misc_feature
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<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43996227

<400> 5368
tgtgtcaggc cagggagtcct ctggctgccc tggatagagt ggagggccct

50

<210> 5369
<211> 50
<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (5370 is other entry)

<221> misc_feature
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<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43996450

<400> 5369
attaagaaag aagaaagaaa aaaaacaacc aaaaacctgg agaataaaca

50

<210> 5370
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5369 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43996450

<400> 5370
attaagaaag aagaaagaaa aaaaaacaac caaaacctg gagaataaac a 51

<210> 5371
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5372 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43996554

<400> 5371
aaggaagaaa gaaaagaaaa agaagaaaa aacggaagca caaatcttcc a 51

<210> 5372
<211> 50
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5371 is other entry)

<221> misc_feature
<222> (25)...(26)
<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43996554

<400> 5372
aaggaagaaa gaaaagaaaa agaagaaaa acggaagcac aaatcttcca 50

<210> 5373
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5374 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43996590

<400> 5373
gagcccggcc ttacctggtg tagggctgtg agtcgcctcct cattgcacaa a 51

<210> 5374
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants. (5373 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43996590

<400> 5374
gagcccgcc ttacctggtg taggggtgtg agtccgtcct cattgcacaa a 51

<210> 5375
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5376 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43996743

<400> 5375
agcaaggga cagagtcacc aaaacctgtc ccaacaatcc ccagacaata t 51

<210> 5376
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5375 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43996743

<400> 5376
agcaaggga cagagtcacc aaaacttgtc ccaacaatcc ccagacaata t 51

<210> 5377
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5378 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43996869

<400> 5377
tctatcataa gaatgtgaac gaggctgaag atctctggac caagatcttg a 51

<210> 5378
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (5377 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43996869

<400> 5378
tctatcataa gaatgtgaac gaggccgaag atctctggac caagatcttg a 51

<210> 5379
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5380 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43997723

<400> 5379
gcgcgccgcc gccacggag gacccactc caacgtgtaa acagaaacag a 51

<210> 5380
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (5379 is other entry)

<221> misc_feature

<222> (0)...(0)
<223> Accession number cg43997723

<400> 5380
gcgcgcgcgc gccacggag gaccctactc caacgtgtaa acagaaacag a 51

<210> 5381
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5382 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43997723

<400> 5381
tggccctgc gtggccagca gggccggctc ctccggaggc tccctgcctt g 51

<210> 5382
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (5381 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43997723

<400> 5382
tggccctgc gtggccagca gggccagctc ctccggaggc tccctgcctt g 51

<210> 5383
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5384 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43997865

<400> 5383
ccggctccag caagagttcc agccactccc ccggctcctg ggtgtgcacc a 51

<210> 5384
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (5383 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43997865

<400> 5384
ccggctccag caagagttcc agccattccc ccggctcctg ggtgtgcacc a 51

<210> 5385
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5386 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43997865

<400> 5385
gtagcggcgg cggcgcagct cgggctccag ttctcggctc agcgtctcct c 51

<210> 5386
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5385 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43997865

<400> 5386
gtagcggcgg cggcgcagct cgggccccag ttctcggctc agcgtctcct c 51

<210> 5387
<211> 51
<212> DNA
<213> Homo sapiens

<220>

<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5388 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43998250

<400> 5387
gagaagtccc cacttaaaaa aaaaaatatc tgcagtttga agggcaaagg g 51

<210> 5388
<211> 50
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (5387 is other entry)

<221> misc_feature
<222> (25)...(26)
<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43998250

<400> 5388
gagaagtccc cacttaaaaa aaaaatatct gcagtttgaa gggcaaaggg 50

<210> 5389
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5390 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43998355

<400> 5389
caccactaca ccttgagggg aacatgcagt cattttcccg gtacagtctg t 51

<210> 5390
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)

<223> 2 of 2 allelic variants (5389 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43998355

<400> 5390

caccactaca ccttgagggg aacatacagt cattttcccg gtacagtctg t

51

<210> 5391

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (5392 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43998922

<400> 5391

aggagggagg aaaagctcct aatgggagag aggggtgcag taagagttta c

51

<210> 5392

<211> 50

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (5391 is other entry)

<221> misc_feature

<222> (25)...(26)

<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43998922

<400> 5392

aggagggagg aaaagctcct aatggagaga aggggtgcagt aagagtttac

50

<210> 5393

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (5394 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43998972

<400> 5393
gtttggagga tcattccaggt cacacgtctc aggtagaaat gcagggtttg a 51

<210> 5394
<211> 50
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (5393 is other entry)

<221> misc_feature
<222> (25)...(26)
<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43998972

<400> 5394
gtttggagga tcattccaggt cacactctca ggtagaaatg cagggtttga 50

<210> 5395
<211> 51
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<220>
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<223> 1 of 2 allelic variants (5396 is other entry)

<221> misc_feature
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<223> Accession number cg43999061

<400> 5395
agcagatagt tgtcagaagt gcataaaaaa actggctttc aggtcatata t 51

<210> 5396
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (5395 is other entry)

<221> misc_feature
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<223> Accession number cg43999061

<400> 5396

agcagatagt tgtcagaagt gcatacaaaa actggctttc aggtcatata t

51

<210> 5397

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 1 of 2 allelic variants (5398 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43999507

<400> 5397

attctagaaa gcaaataaag tcactctcta caataaatag agcatcatgt g

51

<210> 5398

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (5397 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43999507

<400> 5398

attctagaaa gcaaataaag tcactttcta caataaatag agcatcatgt g

51

<210> 5399

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 1 of 2 allelic variants (5400 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43999507

<400> 5399

gagcatcatg tgcttcacag cagacgcgac agagacacat aggccccgtg c

51

<210> 5400

<211> 51
<212> DNA
<213> Homo sapiens

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<223> 2 of 2 allelic variants (5399 is other entry)

<221> misc_feature
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<223> Accession number cg43999507

<400> 5400
gagcatcatg tgcttcacag cagacccgac agagacacat aggccccgtg c 51

<210> 5401
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (5402 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43999507

<400> 5401
ccgtgccaca gcaggatctg agccgtttcc cgttctaaaa gagcatttta a 51

<210> 5402
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (5401 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43999507

<400> 5402
ccgtgccaca gcaggatctg agccgtttcc cgttctaaaa gagcatttta a 51

<210> 5403
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5404 is other entry)

<221> misc_feature
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<223> Accession number cg43999507

<400> 5403
ccacagcagg atctgagccg tttcccggttc taaaagagca ttttaaaaaa t 51

<210> 5404
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (5403 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43999507

<400> 5404
ccacagcagg atctgagccg tttcctgttc taaaagagca ttttaaaaaa t 51

<210> 5405
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (5406 is other entry)

<221> misc_feature
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<223> Accession number cg43999507

<400> 5405
agccgtttcc cgttctaaaa gagcatttta aaaaatgaga ctgagagaga g 51

<210> 5406
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (5405 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43999507

<400> 5406
agccgtttcc cgttctaaaa gagcagttta aaaaatgaga ctacagagaga g 51

<210> 5407
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5408 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43999702

<400> 5407
agtttggcct cgttctcccg ctcccgctgc tcatcgaggt ggctggctg g 51

<210> 5408
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5407 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43999702

<400> 5408
agtttggcct cgttctcccg ctccactgc tcatcgaggt ggctggctg g 51

<210> 5409
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5410 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43999702

<400> 5409
atgcattgtg gacggttctc caaataaaaa aagccccaag ggtttgtcta c 51

<210> 5410
<211> 50

<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5409 is other entry)

<221> misc_feature
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<221> misc_feature
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<223> Accession number cg43999702

<400> 5410
atgcatgctg gacggttctc caaataaaaa agccccaagg gtttgtctac 50

<210> 5411
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5412 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43999766

<400> 5411
caggctttac caagaccttg gttaagtccc agtcacattt actttctgtc t 51

<210> 5412
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5411 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43999766

<400> 5412
caggctttac caagaccttg gttaaattccc agtcacattt actttctgtc t 51

<210> 5413
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5414 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43999810

<400> 5413
catgaagatg tggacttttt ttttttgaga caagagcctc accctgtcgc c 51

<210> 5414
<211> 50
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5413 is other entry)

<221> misc_feature
<222> (25)...(26)
<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43999810

<400> 5414
catgaagatg tggacttttt tttttgagac aagagcctca cctgtcgcc 50

<210> 5415
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5416 is other entry)

<221> misc_feature
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<223> Accession number cg43999946

<400> 5415
aaggtaccac agcaggttat ggttcataca ggactttaaa tgacccatgt t 51

<210> 5416
<211> 51
<212> DNA
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<220>

<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (5415 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43999946

<400> 5416
aaggtaccac agcagggttat ggttcgtaca ggactttaaa tgacccatgt t 51

<210> 5417
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5418 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43999946

<400> 5417
tgacccatgt tgacaataca atttgcaaaa aatatgagaa aagcaataca t 51

<210> 5418
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (5417 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43999946

<400> 5418
tgacccatgt tgacaataca atttgtaaaa aatatgagaa aagcaataca t 51

<210> 5419
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5420 is other entry)

<221> misc_feature
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<223> Accession number cg43999946

<400> 5419

aatatgagaa aagcaatata tatttctgaa acaaaaacat acctgttgta a

51

<210> 5420

<211> 51

<212> DNA

<213> Homo sapiens

<220>

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<222> (26)...(0)

<223> 2 of 2 allelic variants (5419 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43999946

<400> 5420

aatatgagaa aagcaatata tatttttgaa acaaaaacat acctgttgta a

51

<210> 5421

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (5422 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43999946

<400> 5421

cagaatagaa cagttaacag agcagcagtt actgaagaca gacggcaaca g

51

<210> 5422

<211> 51

<212> DNA

<213> Homo sapiens

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<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (5421 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43999946

<400> 5422

cagaatagaa cagttaacag agcagtagtt actgaagaca gacggcaaca g

51

<210> 5423

<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5424 is other entry)

<221> misc_feature
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<223> Accession number cg43999946

<400> 5423
cagagcagca gttactgaag acagacggca acaggcagct ggcgcagcca t 51

<210> 5424
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5423 is other entry)

<221> misc_feature
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<223> Accession number cg43999946

<400> 5424
cagagcagca gttactgaag acagatggca acaggcagct ggcgcagcca t 51

<210> 5425
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5426 is other entry)

<221> misc_feature
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<223> Accession number cg44000241

<400> 5425
ctccctcacg gagccagcgg ccgggaatgc agacatcaga acgtgagggg a 51

<210> 5426
<211> 50
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature

<222> (26)...(0)
<223> 2 of 2 allelic variants (5425 is other entry)

<221> misc_feature
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<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44000241

<400> 5426
ctccctcacg gagccagcgg ccgggatgca gacatcagaa cgtgagggga 50

<210> 5427
<211> 50
<212> DNA
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5428 is other entry)

<221> misc_feature
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<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44000319

<400> 5427
aacttgaaag acagattaaa aaaaactttt tggcaataat ttagaataat 50

<210> 5428
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5427 is other entry)

<221> misc_feature
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<223> Accession number cg44000319

<400> 5428
aacttgaaag acagattaaa aaaaactttt ttggcaataa tttagaataa t 51

<210> 5429
<211> 51
<212> DNA
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<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5430 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44000319

<400> 5429
tgcagggagc tcttctaggt acttagctgt ttttaagggtc tgcactttac c 51

<210> 5430
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5429 is other entry)

<221> misc_feature
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<223> Accession number cg44000319

<400> 5430
tgcagggagc tcttctaggt acttaactgt ttttaagggtc tgcactttac c 51

<210> 5431
<211> 51
<212> DNA
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<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5432 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44000319

<400> 5431
aacataatta caattctgat tatagcacag aaccagagat ggcaaactga c 51

<210> 5432
<211> 50
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5431 is other entry)

<221> misc_feature

<222> (25)...(26)
<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44000319

<400> 5432
aacataatta caattctgat tatagacaga accagagatg gcaaactgac 50

<210> 5433
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5434 is other entry)

<221> misc_feature
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<223> Accession number cg44000319

<400> 5433
atagcacaga accagagatg gcaaactgac gacccaaaag cagagcagga a 51

<210> 5434
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5433 is other entry)

<221> misc_feature
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<223> Accession number cg44000319

<400> 5434
atagcacaga accagagatg gcaaattgac gacccaaaag cagagcagga a 51

<210> 5435
<211> 51
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<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5436 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44000399

<400> 5435
gccttcaatg atctcacttg ctttcccagg tcgctgagag tcccaccatg t 51

<210> 5436
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5435 is other entry)

<221> misc_feature
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<223> Accession number cg44000399

<400> 5436
gccttcaatg atctcacttg ctttctcagg tcgctgagag tcccaccatg t 51

<210> 5437
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5438 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44000405

<400> 5437
cccttctaata ctgaggaaac taagcgtgaa agaatgtgag catgcataaa a 51

<210> 5438
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5437 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44000405

<400> 5438
cccttctaata ctgaggaaac taagcatgaa agaatgtgag catgcataaa a 51

<210> 5439
<211> 51

<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5440 is other entry)

<221> misc_feature
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<223> Accession number cg44000405

<400> 5439
gagtgttggt tttaacctct gtacaatatt tagaccagc aaatgcagaa a 51

<210> 5440
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (5439 is other entry)

<221> misc_feature
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<223> Accession number cg44000405

<400> 5440
gagtgttggt tttaacctct gtacagtatt tagaccagc aaatgcagaa a 51

<210> 5441
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5442 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44000551

<400> 5441
ccagagggtg caatagagag acaaagaatg ctctcttccc tagatgcaa g 51

<210> 5442
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)

<223> 2 of 2 allelic variants (5441 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44000551

<400> 5442

ccagaggttg caatagagag acaaaaaatg ctctcttccc tagatgccaa g

51

<210> 5443

<211> 51

<212> DNA

<213> Homo sapiens

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<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (5444 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44000551

<400> 5443

tgtcagccta ggaatgaaga accaagaaaa gaatcaaact gagtgccaca a

51

<210> 5444

<211> 51

<212> DNA

<213> Homo sapiens

<220>

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<222> (26)...(0)

<223> 2 of 2 allelic variants (5443 is other entry)

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<222> (0)...(0)

<223> Accession number cg44000551

<400> 5444

tgtcagccta ggaatgaaga accaaaaaaa gaatcaaact gagtgccaca a

51

<210> 5445

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (5446 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44001055

<400> 5445
gttgactgaa ctctggaaag agaaacagca gctgatcctg actgcttttg t 51

<210> 5446
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5445 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44001055

<400> 5446
gttgactgaa ctctggaaag agaaatagca gctgatcctg actgcttttg t 51

<210> 5447
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5448 is other entry)

<221> misc_feature
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<223> Accession number cg44001055

<400> 5447
gatcctgact gcttttgtct gtcatatctc cagaactctg gactgtctgg t 51

<210> 5448
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (5447 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44001055

<400> 5448
gatcctgact gcttttgtct gtcattgttc cagaactctg gactgtctgg t 51

<210> 5449
<211> 51
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<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (5450 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44001055

<400> 5449

ctgcttttgt ctgtcatatt tccagaactc tggactgtct ggttgaattg t

51

<210> 5450

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 2 of 2 allelic variants (5449 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44001055

<400> 5450

ctgcttttgt ctgtcatatt tccaggactc tggactgtct ggttgaattg t

51

<210> 5451

<211> 51

<212> DNA

<213> Homo sapiens

<220>

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<222> (26)...(0)

<223> 1 of 2 allelic variants (5452 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44001055

<400> 5451

tgcttttgtc tgtcatattt ccagaactct ggactgtctg gttgaattgt c

51

<210> 5452

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (5451 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44001055

<400> 5452
tgcttttgtc tgtcatattt ccagagctct ggactgtctg gttgaattgt c 51

<210> 5453
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (5454 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44001055

<400> 5453
cagaactctg gactgtctgg ttgaattgtc ggaggatctc tcgttgcttt c 51

<210> 5454
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (5453 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44001055

<400> 5454
cagaactctg gactgtctgg ttgaactgtc ggaggatctc tcgttgcttt c 51

<210> 5455
<211> 51
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<223> 1 of 2 allelic variants (5456 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44001055

<400> 5455

tctggttgaa ttgtcggagg atctctcggt gctttctgac gtaccaggtg t

51

<210> 5456

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 2 of 2 allelic variants (5455 is other entry)

<221> misc_feature

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<223> Accession number cg44001055

<400> 5456

tctggttgaa ttgtcggagg atctcccggt gctttctgac gtaccaggtg t

51

<210> 5457

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 1 of 2 allelic variants (5458 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44001055

<400> 5457

gagcggcacg cttctgggtc ttcatagggg tgcccttggt gagatgctgg g

51

<210> 5458

<211> 51

<212> DNA

<213> Homo sapiens

<220>

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<222> (26)...(0)

<223> 2 of 2 allelic variants (5457 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44001055

<400> 5458

gagcggcacg cttctgggtc ttcatagggg tgcccttggt gagatgctgg g

51

<210> 5459

<211> 51

<212> DNA

<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (5460 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44001076

<400> 5459
caaagcctgt agttgctgct tcttggttgg aagattctgg acagcctgaa a 51

<210> 5460
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5459 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44001076

<400> 5460
caaagcctgt agttgctgct tcttgattgg aagattctgg acagcctgaa a 51

<210> 5461
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5462 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44001185

<400> 5461
tattggggga tgtcagcaga gaacgtggga catgaaaaca agtcttagga g 51

<210> 5462
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5461 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44001185

<400> 5462
tattggggga tgtcagcaga gaacgcggga catgaaaaca agtcttagga g 51

<210> 5463
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5464 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44001185

<400> 5463
caagacccaa gccttgaccc taagtaacag atgcaaggat cacgaacaac c 51

<210> 5464
<211> 50
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (5463 is other entry)

<221> misc_feature
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<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44001185

<400> 5464
caagacccaa gccttgaccc taagtacaga tgcaaggatc acgaacaacc 50

<210> 5465
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5466 is other entry)

<221> misc_feature
<222> (0)...(0)

<223> Accession number cg44001210

<400> 5465

tgggtatggg gccgttgcc cggggcccg atagtgtgga gggttctctt g

51

<210> 5466

<211> 51

<212> DNA

<213> Homo sapiens

<220>

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<222> (26)...(0)

<223> 2 of 2 allelic variants (5465 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44001210

<400> 5466

tgggtatggg gccgttgcc cggggtcgg atagtgtgga gggttctctt g

51

<210> 5467

<211> 51

<212> DNA

<213> Homo sapiens

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<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (5468 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44001481

<400> 5467

acctgatgaa attattttta aaaagtttat attccacaga gtttcagttt c

51

<210> 5468

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (5467 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44001481

<400> 5468

acctgatgaa attattttta aaaagatttat attccacaga gtttcagttt c

51

<210> 5469

<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5470 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44001539

<400> 5469
gagcgggtcg gccgtgcgga ccccgctgaa gaacagcata acagccgagt c 51

<210> 5470
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5469 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44001539

<400> 5470
gagcgggtcg gccgtgcgga ccccgatgaa gaacagcata acagccgagt c 51

<210> 5471
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5472 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44001588

<400> 5471
aacaataaca acaacgacga gggtcgctgt gaagctgaaa gttgcagagt g 51

<210> 5472
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature

<222> (26)...(0)
<223> 2 of 2 allelic variants (5471 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44001588

<400> 5472
aacaataaca acaacgacga gggctctctgt gaagctgaaa gttgcagagt g 51

<210> 5473
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5474 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44001596

<400> 5473
caatcccagc atagcacaca ccacatata gacgatccat tatcatcgca a 51

<210> 5474
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (5473 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44001596

<400> 5474
caatcccagc atagcacaca ccaccgtata gacgatccat tatcatcgca a 51

<210> 5475
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5476 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44002106

<400> 5475
ccccgggaag agacggcatg aattctaaga gtgccaggg tctggctggt c 51

<210> 5476
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (5475 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44002106

<400> 5476
ccccgggaag agacggcatg aattcaaaga gtgccaggg tctggctggt c 51

<210> 5477
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5478 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44002106

<400> 5477
acacgtgctt catgaactca attctgcagt gcctgagcaa caccgggag t 51

<210> 5478
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5477 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44002106

<400> 5478
acacgtgctt catgaactca attcttcagt gcctgagcaa caccgggag t 51

<210> 5479
<211> 51

<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (5480 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44002106

<400> 5479
gcttcatgaa ctcaattctg cagtgcctga gcaacacccg ggagttgaga g 51

<210> 5480
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5479 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44002106

<400> 5480
gcttcatgaa ctcaattctg cagtgtctga gcaacacccg ggagttgaga g 51

<210> 5481
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5482 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44002106

<400> 5481
ttctgcagtg cctgagcaac acccgaggagt tgagagatta ctgcctccag a 51

<210> 5482
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)

<223> 2 of 2 allelic variants (5481 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44002106

<400> 5482

ttctgcagtg cctgagcaac acccgcgagt tgagagatta ctgcctccag a

51

<210> 5483

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (5484 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44002106

<400> 5483

tccagaggct gtacatgcgg gacctggacc acagcagcag tgcacacaca g

51

<210> 5484

<211> 51

<212> DNA

<213> Homo sapiens

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<222> (26)...(0)

<223> 2 of 2 allelic variants (5483 is other entry)

<221> misc_feature

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<223> Accession number cg44002106

<400> 5484

tccagaggct gtacatgcgg gacctcgacc acagcagcag tgcacacaca g

51

<210> 5485

<211> 51

<212> DNA

<213> Homo sapiens

<220>

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<222> (26)...(0)

<223> 1 of 2 allelic variants (5486 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44002106

<400> 5485
accacagcag cagtgcacac acagccctcg tggaagagtt tgcaaaaacta a 51

<210> 5486
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (5485 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44002106

<400> 5486
accacagcag cagtgcacac acagctctcg tggaagagtt tgcaaaaacta a 51

<210> 5487
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5488 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44002106

<400> 5487
cagcagcagt gcacacacag ccctcgtgga agagtttgca aaactaatcc a 51

<210> 5488
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (5487 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44002106

<400> 5488
cagcagcagt gcacacacag ccctcatgga agagtttgca aaactaatcc a 51

<210> 5489
<211> 51
<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (5490 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44002125

<400> 5489

ggccgttctc ctggaggagg attctcgcca cctgctggcc atgaagctgg a

51

<210> 5490

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (5489 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44002125

<400> 5490

ggccgttctc ctggaggagg attcttgcca cctgctggcc atgaagctgg a

51

<210> 5491

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (5492 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44002131

<400> 5491

actccagagg cagagtggga cctgggctcc cgagggcaga ccgcagcaga g

51

<210> 5492

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (5491 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44002131

<400> 5492
actccagagg cagagtggga cctggcctcc cgagggcaga ccgcagcaga g 51

<210> 5493
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5494 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44002131

<400> 5493
gaggcagagt gggacctggg ctcccagagg cagaccgcag cagagacccc a 51

<210> 5494
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (5493 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44002131

<400> 5494
gaggcagagt gggacctggg ctcccagagg cagaccgcag cagagacccc a 51

<210> 5495
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5496 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44002258

<400> 5495

gcaagtaaca ctgaatgtcc aaaaatacgg ctgtgttaaa ctaacaagcc a

51

<210> 5496

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (5495 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44002258

<400> 5496

gcaagtaaca ctgaatgtcc aaaaacacgg ctgtgttaaa ctaacaagcc a

51

<210> 5497

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (5498 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44002457

<400> 5497

gaggaagtgt cagcaaacat gaaaaagcaa atccggccat aactatatag a

51

<210> 5498

<211> 50

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (5497 is other entry)

<221> misc_feature

<222> (25)...(26)

<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44002457

<400> 5498

gaggaagtgt cagcaaacat gaaaagcaaa tccggccata actatataga

50

<210> 5499
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5500 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44002457

<400> 5499
ttcagccacg tacccgtaga taaatccaac tattgcagaa aaaagaataa t 51

<210> 5500
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5499 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44002457

<400> 5500
ttcagccacg tacccgtaga taaattcaac tattgcagaa aaaagaataa t 51

<210> 5501
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5502 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44002457

<400> 5501
ccatctgcgt gggcagcgag ctcagatgct ccagcatggc tggctgagga g 51

<210> 5502
<211> 51
<212> DNA
<213> Homo sapiens

<220>

<221> misc_feature
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<223> 2 of 2 allelic variants (5501 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44002457

<400> 5502
ccatctgcgt gggcagcgag ctcaggtgct ccagcatggc tggctgagga g 51

<210> 5503
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5504 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44002584

<400> 5503
ttagcgtagg ctgctgctgc tgctggcctg ggagctgccc atacctgggt g 51

<210> 5504
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (5503 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44002584

<400> 5504
ttagcgtagg ctgctgctgc tgctgacctg ggagctgccc atacctgggt g 51

<210> 5505
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5506 is other entry)

<221> misc_feature
<222> (0)...(0)

<223> Accession number cg44002584

<400> 5505

tcaggcagga agttacttag cttctccttc accttcttcc cacagaattt a

51

<210> 5506

<211> 51

<212> DNA

<213> Homo sapiens

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<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (5505 is other entry)

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<222> (0)...(0)

<223> Accession number cg44002584

<400> 5506

tcaggcagga agttacttag cttctccttc accttcttcc cacagaattt a

51

<210> 5507

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 1 of 2 allelic variants (5508 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44002588

<400> 5507

tctgtctatg gcaataatac gatgcatatt gagaaactgc ttcaaagatg g

51

<210> 5508

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 2 of 2 allelic variants (5507 is other entry)

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<223> Accession number cg44002588

<400> 5508

tctgtctatg gcaataatac gatgcatatt gagaaactgc ttcaaagatg g

51

<210> 5509

<211> 51
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<223> 1 of 2 allelic variants (5510 is other entry)

<221> misc_feature
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<223> Accession number cg44002594

<400> 5509
cgccctggccg ttggcgctgc ccagccgcct gcagatcagg ctctggatgc c 51

<210> 5510
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (5509 is other entry)

<221> misc_feature
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<223> Accession number cg44002594

<400> 5510
cgccctggccg ttggcgctgc ccagctgcct gcagatcagg ctctggatgc c 51

<210> 5511
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (5512 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44002594

<400> 5511
ccccatccaa ggaattgtgg catctcggag ggctggaccg gcacacgccg g 51

<210> 5512
<211> 51
<212> DNA
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5511 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44002594

<400> 5512
ccccatccaa ggaattgtgg catcttggag ggctggaccg gcacacgccg g 51

<210> 5513
<211> 51
<212> DNA
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<220>
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<223> 1 of 2 allelic variants (5514 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44003017

<400> 5513
catctattga atctttggct ttgtcgttgc aatgcatggt gcaccgggcc a 51

<210> 5514
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (5513 is other entry)

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<223> Accession number cg44003017

<400> 5514
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<210> 5515
<211> 51
<212> DNA
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<220>
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<223> 1 of 2 allelic variants (5516 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44003017

<400> 5515
gttgcaatgc atggtgcacc gggccaggcg gtcttgaac ttctccagct c 51

<210> 5516
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (5515 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44003017

<400> 5516
gttgcaatgc atggtgcacc gggccgggcg gtcttgaac ttctccagct c 51

<210> 5517
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (5518 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44003792

<400> 5517
tgccagggaa ggaggaccct ataggtggc cagcaagggg ccactggcgg t 51

<210> 5518
<211> 50
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (5517 is other entry)

<221> misc_feature
<222> (25)...(26)
<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44003792

<400> 5518

tgccagggaa ggaggaccct ataggtggcc agcaaggggc cactggcggt 50

<210> 5519

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 1 of 2 allelic variants (5520 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44003959

<400> 5519

gccccctcc actctgcagg gcctccacac gcacacgggg ccggcgacgg a 51

<210> 5520

<211> 51

<212> DNA

<213> Homo sapiens

<220>

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<222> (26)...(0)

<223> 2 of 2 allelic variants (5519 is other entry)

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<222> (0)...(0)

<223> Accession number cg44003959

<400> 5520

gccccctcc actctgcagg gcctcaacac gcacacgggg ccggcgacgg a 51

<210> 5521

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (5522 is other entry)

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<400> 5521

tcaccatggt gccagactg gtcttgaact cctgggtca agcaatccat c 51

<210> 5522

<211> 51

<212> DNA

<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (5521 is other entry)

<221> misc_feature
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<223> Accession number cg44004084

<400> 5522
tcaccatggt gccagactg gtcttaaact cctgggctca agcaatccat c 51

<210> 5523
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<220>
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<223> 1 of 2 allelic variants (5524 is other entry)

<221> misc_feature
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<223> Accession number cg44004084

<400> 5523
gccagactg gtcttgaact cctgggctca agcaatccat ccaccttggc c 51

<210> 5524
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5523 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44004084

<400> 5524
gccagactg gtcttgaact cctggcctca agcaatccat ccaccttggc c 51

<210> 5525
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5526 is other entry)

<221> misc_feature
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<223> Accession number cg44004084

<400> 5525
ttgaactcct gggctcaagc aatccatcca ccttggcctc ccaaagtgct g 51

<210> 5526
<211> 51
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<220>
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<223> 2 of 2 allelic variants (5525 is other entry)

<221> misc_feature
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<223> Accession number cg44004084

<400> 5526
ttgaactcct gggctcaagc aatccttcca ccttggcctc ccaaagtgct g 51

<210> 5527
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5528 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44004120

<400> 5527
ggacttccta tgtcactttc caagggttc agccacttg aaggccaag g 51

<210> 5528
<211> 50
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5527 is other entry)

<221> misc_feature
<222> (25)...(26)
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<221> misc_feature
<222> (0)...(0)

<223> Accession number cg44004120

<400> 5528

ggacttccta tgtcactttc caaggcttca gccacttga agggccaagg

50

<210> 5529

<211> 51

<212> DNA

<213> Homo sapiens

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<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (5530 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44004303

<400> 5529

agggtgtagt ccacccgccca gccacacct gccaacctat tcatgcgtag g

51

<210> 5530

<211> 50

<212> DNA

<213> Homo sapiens

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<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (5529 is other entry)

<221> misc_feature

<222> (25)...(26)

<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44004303

<400> 5530

agggtgtagt ccacccgccca gcccaacctg ccaacctatt catgcgtagg

50

<210> 5531

<211> 51

<212> DNA

<213> Homo sapiens

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<222> (26)...(0)

<223> 1 of 2 allelic variants (5532 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44004427

<400> 5531
atccctgaat taacaagcag ccagtggcga gtgttcacga taaaactgaa t 51

<210> 5532
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (5531 is other entry)

<221> misc_feature
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<223> Accession number cg44004427

<400> 5532
atccctgaat taacaagcag ccagtcgcga gtgttcacga taaaactgaa t 51

<210> 5533
<211> 51
<212> DNA
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<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5534 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44004474

<400> 5533
ccgaggattg agagctccca atattctttg gagaataagc agtagttttg c 51

<210> 5534
<211> 50
<212> DNA
<213> Homo sapiens

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<223> 2 of 2 allelic variants (5533 is other entry)

<221> misc_feature
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<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44004474

<400> 5534
ccgaggattg agagctccca atattttttg agaataagca gtagttttgc 50

<210> 5535
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5536 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44004688

<400> 5535
gcctgtcagc tgggtgggcag ccctggagtg tggatggaag aacaggcatg c 51

<210> 5536
<211> 51
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<220>
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<223> 2 of 2 allelic variants (5535 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44004688

<400> 5536
gcctgtcagc tgggtgggcag ccctgaagtg tggatggaag aacaggcatg c 51

<210> 5537
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 1 of 2 allelic variants (5538 is other entry)

<221> misc_feature
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<223> Accession number cg44004731

<400> 5537
ctgtgggtct tcttttttct gtggcaaatt caccttctca aaaacaacag g 51

<210> 5538
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5537 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44004731

<400> 5538
ctgtgggtct tcttttttct gtggctaatt caccttctca aaaacaacag g 51

<210> 5539
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5540 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44004731

<400> 5539
tgtgggtctt ctttttttctg tggcaaattc accttctcaa aaacaacagg t 51

<210> 5540
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5539 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44004731

<400> 5540
tgtgggtctt ctttttttctg tggcatattc accttctcaa aaacaacagg t 51

<210> 5541
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5542 is other entry)

<221> misc_feature

<222> (0)...(0)
<223> Accession number cg44004731

<400> 5541
gtgggtcttc ttttttctgt ggcaaattca ccttctcaaa aacaacaggt t 51

<210> 5542
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5541 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44004731

<400> 5542
gtgggtcttc ttttttctgt ggcaatttca ccttctcaaa aacaacaggt t 51

<210> 5543
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5544 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44004731

<400> 5543
aagttttctt gtctgaattt tcaagtgggg tgaacaatga ctgagaggaa a 51

<210> 5544
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5543 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44004731

<400> 5544
aagttttctt gtctgaattt tcaagggggg tgaacaatga ctgagaggaa a 51

<210> 5545
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5546 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44004823

<400> 5545
gctgccacag tcctgaccct ggccctgctg ggcaatgcc atgctgctc c

51

<210> 5546
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5545 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44004823

<400> 5546
gctgccacag tcctgaccct ggccccgctg ggcaatgcc atgctgctc c

51

<210> 5547
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5548 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44004831

<400> 5547
ggactcacc aacaaaatgt gctctgttaa cacaaccagc agtacaatca t

51

<210> 5548
<211> 51
<212> DNA
<213> Homo sapiens

<220>

<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (5547 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44004831

<400> 5548
ggactcaccc aacaaaatgt gctctattaa cacaaccagc agtacaatca t 51

<210> 5549
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5550 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44005101

<400> 5549
agtgcctttt gaggcaaatc catacgtcgt cggggagcaa aggattgctg a 51

<210> 5550
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (5549 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44005101

<400> 5550
agtgcctttt gaggcaaatc catacctcgt cggggagcaa aggattgctg a 51

<210> 5551
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5552 is other entry)

<221> misc_feature
<222> (0)...(0)

<223> Accession number cg44005101

<400> 5551
gccctttgag gcaaattccat acgtcgtcgg ggagcaaagg attgctgac t 51

<210> 5552
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5551 is other entry)

<221> misc_feature
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<223> Accession number cg44005101

<400> 5552
gccctttgag gcaaattccat acgtcctcgg ggagcaaagg attgctgac t 51

<210> 5553
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5554 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44005101

<400> 5553
ctttgaggca aatccatcag tcgtcgggga gcaaaggatt gctgatctct g 51

<210> 5554
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5553 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44005101

<400> 5554
ctttgaggca aatccatcag tcgtcaggga gcaaaggatt gctgatctct g 51

<210> 5555

<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5556 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44005150

<400> 5555
acactgaaat cagagcctgc acacagagca gcagatgctt caatgtaaag g

51

<210> 5556
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (5555 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44005150

<400> 5556
acactgaaat cagagcctgc acacaaagca gcagatgctt caatgtaaag g

51

<210> 5557
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5558 is other entry)

<221> misc_feature
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<223> Accession number cg44005150

<400> 5557
acacagagca gcagatgctt caatgtaaag gtcatttcca ggtccttgac a

51

<210> 5558
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature

<222> (26)...(0)
<223> 2 of 2 allelic variants (5557 is other entry)

<221> misc_feature
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<223> Accession number cg44005150

<400> 5558
acacagagca gcagatgctt caatgcaaag gtcatttcca ggtccttgac a 51

<210> 5559
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5560 is other entry)

<221> misc_feature
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<223> Accession number cg44005468

<400> 5559
ccaggagtca cggatgggaa agtaaattctt tggaggggct gggagctggg g 51

<210> 5560
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (5559 is other entry)

<221> misc_feature
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<223> Accession number cg44005468

<400> 5560
ccaggagtca cggatgggaa agtaagtctt tggaggggct gggagctggg g 51

<210> 5561
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5562 is other entry)

<221> misc_feature
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<223> Accession number cg44005989

<400> 5561
tcttgccgc agactgagcc tgtacctcac ccgtctccca ccaactcttg g 51

<210> 5562
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5561 is other entry)

<221> misc_feature
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<223> Accession number cg44005989

<400> 5562
tcttgccgc agactgagcc tgtacttcac ccgtctccca ccaactcttg g 51

<210> 5563
<211> 51
<212> DNA
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<223> 1 of 2 allelic variants (5564 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44006130

<400> 5563
aatggcagca cttccgagag tggcacagag tctgttggtg gccagcacag g 51

<210> 5564
<211> 51
<212> DNA
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<223> 2 of 2 allelic variants (5563 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44006130

<400> 5564
aatggcagca cttccgagag tggcatagag tctgttggtg gccagcacag g 51

<210> 5565
<211> 51

<212> DNA
<213> Homo sapiens

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<223> 1 of 2 allelic variants (5566 is other entry)

<221> misc_feature
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<223> Accession number cg44006486

<400> 5565
ctccctcact cattcattat tcacacagag tatataacag tttttttttt t 51

<210> 5566
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 2 of 2 allelic variants (5565 is other entry)

<221> misc_feature
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<223> Accession number cg44006486

<400> 5566
ctccctcact cattcattat tcacaaagag tatataacag tttttttttt t 51

<210> 5567
<211> 51
<212> DNA
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<220>
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<223> 1 of 2 allelic variants (5568 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44006536

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ttatggtttc gactaacatt aagtatacct ttttttgaat caacaggatg a 51

<210> 5568
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 2 of 2 allelic variants (5567 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44006536

<400> 5568

ttatgggttc gactaacatt aagtacacct ttttttgaat caacaggatg a

51

<210> 5569

<211> 51

<212> DNA

<213> Homo sapiens

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<222> (26)...(0)

<223> 1 of 2 allelic variants (5570 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44006660

<400> 5569

aagacctgtg gagaccatca tcgaggccat ggccccacat ctgtgatatg g

51

<210> 5570

<211> 50

<212> DNA

<213> Homo sapiens

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<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44006660

<400> 5570

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50

<210> 5571

<211> 51

<212> DNA

<213> Homo sapiens

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<222> (26)...(0)

<223> 1 of 2 allelic variants (5572 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44007198

<400> 5571
ttataacaac atgatcccca cagtcaactca gtgcatggta cacaggttta t 51

<210> 5572
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 2 of 2 allelic variants (5571 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44007198

<400> 5572
ttataacaac atgatcccca cagtcgctca gtgcatggta cacaggttta t 51

<210> 5573
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5574 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44007371

<400> 5573
gtacagtggg aatccttcca tcatacagggt aatatataat aacattcaaa a 51

<210> 5574
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (5573 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44007371

<400> 5574
gtacagtggg aatccttcca tcatatagggt aatatataat aacattcaaa a 51

<210> 5575
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5576 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44007434

<400> 5575
tatggacccc tgaccccgcg gggtcgttcg gactcttaac gtgtggactg a 51

<210> 5576
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (5575 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44007434

<400> 5576
tatggacccc tgaccccgcg gggtcattcg gactcttaac gtgtggactg a 51

<210> 5577
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5578 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44008138

<400> 5577
tgctgggttta atataattag tataactaaat agttttctgc atttatttgg t 51

<210> 5578
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (5577 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44008138

<400> 5578
tgctggttta atataattag tatacgaaat agttttctgc atttatttg t 51

<210> 5579
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5580 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44008187

<400> 5579
gtgcaggtcc cacacaacag gcaaaagctc tagctttggc ctggaggcag c 51

<210> 5580
<211> 50
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (5579 is other entry)

<221> misc_feature
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<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44008187

<400> 5580
gtgcaggtcc cacacaacag gcaaagctct agctttggcc tggaggcagc 50

<210> 5581
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5582 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44009119

<400> 5581
aggtggtgga gtctgagatt tagaggctga gcctttgggg gtgggggcag a 51

<210> 5582
<211> 50
<212> DNA
<213> Homo sapiens

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<223> 2 of 2 allelic variants (5581 is other entry)

<221> misc_feature
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<221> misc_feature
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<223> Accession number cg44009119

<400> 5582
aggtggtgga gtctgagatt tagagctgag cctttggggg tgggggcaga 50

<210> 5583
<211> 51
<212> DNA
<213> Homo sapiens

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<222> (26)...(0)
<223> 1 of 2 allelic variants (5584 is other entry)

<221> misc_feature
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<223> Accession number cg44009213

<400> 5583
taataccctc ccccatcctt aactctagaa ccccggtttg gtggggagga g 51

<210> 5584
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 2 of 2 allelic variants (5583 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44009213

<400> 5584
taataccctc ccccatcctt aactccagaa ccccggtttg gtggggagga g 51

<210> 5585
<211> 51
<212> DNA
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5586 is other entry)

<221> misc_feature
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<223> Accession number cg44009645

<400> 5585
gcagagtggg ggcaatgtca tgtgggcaca tgcccgctgc tctgctaatt g 51

<210> 5586
<211> 50
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5585 is other entry)

<221> misc_feature
<222> (25)...(26)
<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44009645

<400> 5586
gcagagtggg ggcaatgtca tgtggcacat gcccgctgct ctgctaattg 50

<210> 5587
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (5588 is other entry)

<221> misc_feature

<222> (0)...(0)
<223> Accession number cg44009958

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ccaaaaatac catatgcatt tcaggtgtca ttcattatct ctcaagtcaca a 51

<210> 5588
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5587 is other entry)

<221> misc_feature
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<223> Accession number cg44009958

<400> 5588
ccaaaaatac catatgcatt tcaggcgtca ttcattatct ctcaagtcaca a 51

<210> 5589
<211> 51
<212> DNA
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<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5590 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44010401

<400> 5589
agtctgtttt tgaaacagct ttccacttca tctccctttc tggggctcag g 51

<210> 5590
<211> 51
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<213> Homo sapiens

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<223> 2 of 2 allelic variants (5589 is other entry)

<221> misc_feature
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<223> Accession number cg44010401

<400> 5590
agtctgtttt tgaaacagct ttccatttca tctccctttc tggggctcag g 51

<210> 5591
<211> 51
<212> DNA
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<223> 1 of 2 allelic variants (5592 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44010528

<400> 5591
ggcatggatg atacctctct tcaggcgctc catggcgctc ttgctgccag c 51

<210> 5592
<211> 50
<212> DNA
<213> Homo sapiens

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<222> (26)...(0)
<223> 2 of 2 allelic variants (5591 is other entry)

<221> misc_feature
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<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
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<223> Accession number cg44010528

<400> 5592
ggcatggatg atacctctct tcagggtccc atggcgctct tgctgccagc 50

<210> 5593
<211> 51
<212> DNA
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<220>
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<223> 1 of 2 allelic variants (5594 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44010528

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ctggagccag ggggtgctgct gggagcaggg gcgatggggt ttaggacat c .51

<210> 5594
<211> 51

<212> DNA
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<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5593 is other entry)

<221> misc_feature
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<223> Accession number cg44010528

<400> 5594
ctggagccag gggtgctgct gggagtaggg gcgatggggt ttaggacat c 51

<210> 5595
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5596 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44010626

<400> 5595
gggctggggc tgggccttcc aagacgatcg acagaaccac caccaggacc g 51

<210> 5596
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (5595 is other entry)

<221> misc_feature
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<223> Accession number cg44010626

<400> 5596
gggctggggc tgggccttcc aagaccatcg acagaaccac caccaggacc g 51

<210> 5597
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (5598 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44010683

<400> 5597

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<210> 5598

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 2 of 2 allelic variants (5597 is other entry)

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<223> Accession number cg44010683

<400> 5598

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51

<210> 5599

<211> 51

<212> DNA

<213> Homo sapiens

<220>

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<222> (26)...(0)

<223> 1 of 2 allelic variants (5600 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44011509

<400> 5599

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51

<210> 5600

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 2 of 2 allelic variants (5599 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44011509

<400> 5600
acaagccact ggatctgtcc gattctacat tgtcttacac tgaaacggag g 51

<210> 5601
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5602 is other entry)

<221> misc_feature
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<223> Accession number cg44011509

<400> 5601
attccacatt gtcttacact gaaacggagg ctaccaactc cctcatcact g 51

<210> 5602
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5601 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44011509

<400> 5602
attccacatt gtcttacact gaaacggagg ctaccaactc cctcatcact g 51

<210> 5603
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5604 is other entry)

<221> misc_feature
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<223> Accession number cg44011509

<400> 5603
cattgtctta cactgaaacg gaggtacca actccctcat cactgtccg g 51

<210> 5604
<211> 51
<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (5603 is other entry)

<221> misc_feature

<222> (0)...(0)

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<400> 5604

cattgtctta cactgaaacg gaggccacca actccctcat cactgctccg g

51

<210> 5605

<211> 51

<212> DNA

<213> Homo sapiens

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<222> (26)...(0)

<223> 1 of 2 allelic variants (5606 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44011509

<400> 5605

tcactgctcc gggatgaattc tcagacgccca gcatgtctcc ggacgccacc a

51

<210> 5606

<211> 51

<212> DNA

<213> Homo sapiens

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<222> (26)...(0)

<223> 2 of 2 allelic variants (5605 is other entry)

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<222> (0)...(0)

<223> Accession number cg44011509

<400> 5606

tcactgctcc gggatgaattc tcagatgccca gcatgtctcc ggacgccacc a

51

<210> 5607

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (5608 is other entry)

<221> misc_feature
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<223> Accession number cg44011509

<400> 5607
tctcagacgc cagcatgtct ccggacgccca ccaagccgag ccactggtgc a 51

<210> 5608
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (5607 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44011509

<400> 5608
tctcagacgc cagcatgtct ccggatgccca ccaagccgag ccactggtgc a 51

<210> 5609
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5610 is other entry)

<221> misc_feature
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<223> Accession number cg44011594

<400> 5609
tgtcactcat tcttatcact ctgtctttct tggccaacc catcagctgg c 51

<210> 5610
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (5609 is other entry)

<221> misc_feature
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<223> Accession number cg44011594

<400> 5610

tgtcactcat tcttatcact ctgtccttct tggccaacc catcagctgg c

51

<210> 5611

<211> 51

<212> DNA

<213> Homo sapiens

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<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (5612 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44011736

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ccacgtgggg ctcagacggg cattgtgctg ctctgggagg caagtccatg g

51

<210> 5612

<211> 51

<212> DNA

<213> Homo sapiens

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<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (5611 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44011736

<400> 5612

ccacgtgggg ctcagacggg cattgagctg ctctgggagg caagtccatg g

51

<210> 5613

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (5614 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44012172

<400> 5613

gaacgcggc tcttcgcctc tcagcggcgg cttgtccttt gttccggacg c

51

<210> 5614

<211> 50

<212> DNA

<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5613 is other entry)

<221> misc_feature
<222> (25)...(26)
<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44012172

<400> 5614
gaacgcgcgc tcttcgcctc tcagcgcggc ttgtcctttg ttccggacgc 50

<210> 5615
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5616 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44012500

<400> 5615
gtaactggag agagattcct tcacctgggt gaggaaggtc gggctaggca t 51

<210> 5616
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (5615 is other entry)

<221> misc_feature
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<223> Accession number cg44012500

<400> 5616
gtaactggag agagattcct tcacccgggt gaggaaggtc gggctaggca t 51

<210> 5617
<211> 51
<212> DNA
<213> Homo sapiens

<220>

<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5618 is other entry)

<221> misc_feature
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<223> Accession number cg44012840

<400> 5617
acttttctgt cccaataatt gagagggtg tttcatttcc aaaaaaggga a 51

<210> 5618
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<212> DNA
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<220>
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<221> misc_feature
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<223> Accession number cg44012840

<400> 5618
acttttctgt cccaataatt gagagtgtg tttcatttcc aaaaaaggga a 51

<210> 5619
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5620 is other entry)

<221> misc_feature
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<223> Accession number cg44012840

<400> 5619
cagtgaatct atgaattgtt taagagaaag gtcactccgt tactgacttc t 51

<210> 5620
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<212> DNA
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<220>
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<223> 2 of 2 allelic variants (5619 is other entry)

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<223> Accession number cg44012840

<400> 5620

cagtgaatct atgaattggt taagacaaag gtcactccgt tactgacttc t 51

<210> 5621

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 1 of 2 allelic variants (5622 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44012840

<400> 5621

gttactgact tctgctacat ctaatatcc agggaagtaa tatttagaga t 51

<210> 5622

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 2 of 2 allelic variants (5621 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44012840

<400> 5622

gttactgact tctgctacat ctaattttcc agggaagtaa tatttagaga t 51

<210> 5623

<211> 51

<212> DNA

<213> Homo sapiens

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<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (5624 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44012940

<400> 5623

gtgagcctcc aggattcagg ggttctgggg aggacagatt tgctcggggg t 51

<210> 5624

<211> 51
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<223> 2 of 2 allelic variants (5623 is other entry)

<221> misc_feature
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<400> 5624
gtgagcctcc aggattcagg ggttcgagg aggacagatt tgctcggggt g 51

<210> 5625
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (5626 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44013863

<400> 5625
tggtccgacc gtggatggtg attttctcac tgacatgcc a gacatattac t 51

<210> 5626
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5625 is other entry)

<221> misc_feature
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<223> Accession number cg44013863

<400> 5626
tggtccgacc gtggatggtg attttttcac tgacatgcc a gacatattac t 51

<210> 5627
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature

<222> (26)...(0)
<223> 1 of 2 allelic variants (5628 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44013918

<400> 5627
ttcaaagtct ctatccctat cccagatagg ccacttggcc cagggcaggg c 51

<210> 5628
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 2 of 2 allelic variants (5627 is other entry)

<221> misc_feature
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<223> Accession number cg44013918

<400> 5628
ttcaaagtct ctatccctat cccagttagg ccacttggcc cagggcaggg c 51

<210> 5629
<211> 51
<212> DNA
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<223> 1 of 2 allelic variants (5630 is other entry)

<221> misc_feature
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<223> Accession number cg44014420

<400> 5629
tcctcttcat ttgacccaaa atatcctggg aggtccagca tcctctgctc a 51

<210> 5630
<211> 50
<212> DNA
<213> Homo sapiens

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<223> 2 of 2 allelic variants (5629 is other entry)

<221> misc_feature
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<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
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<223> Accession number cg44014420

<400> 5630
tcctcttcat ttgacccaaa atatctggga ggtccagcat cctctgctca 50

<210> 5631
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (5632 is other entry)

<221> misc_feature
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<223> Accession number cg44014643

<400> 5631
caccacgcgt gtgaggtggg tcccagaggg gggccccaga tggggccagt c 51

<210> 5632
<211> 50
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (5631 is other entry)

<221> misc_feature
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<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
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<223> Accession number cg44014643

<400> 5632
caccacgcgt gtgaggtggg tcccaagggg ggccccagat ggggccagtc 50

<210> 5633
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (5634 is other entry)

<221> misc_feature

<222> (0)...(0)
<223> Accession number cg44014643

<400> 5633
cgtgtgaggt ggggtcccaga ggggggcccc agatggggcc agtcctgtga g 51

<210> 5634
<211> 50
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (5633 is other entry)

<221> misc_feature
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<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
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<223> Accession number cg44014643

<400> 5634
cgtgtgaggt ggggtcccaga gggggcccca gatggggcca gtcctgtgag 50

<210> 5635
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (5636 is other entry)

<221> misc_feature
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<223> Accession number cg44014643

<400> 5635
ggggctggag gtgccccgcg gcatgttttg tgtggcaaca ctgataacct a 51

<210> 5636
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (5635 is other entry)

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<223> Accession number cg44014643

<400> 5636
ggggctggag gtgccccgcg gcatgctttg tgtggcaaca ctgataacct a 51

<210> 5637
<211> 39
<212> DNA
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<220>
<221> misc_feature
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<223> 1 of 2 allelic variants (5638 is other entry)

<221> misc_feature
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<223> Accession number cg44014666

<400> 5637
cggagatctg ctgacgcggt ctacccttcc ggcccgtgt 39

<210> 5638
<211> 39
<212> DNA
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<223> 2 of 2 allelic variants (5637 is other entry)

<221> misc_feature
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<223> Accession number cg44014666

<400> 5638
cggagatctg ctgccgcggt ctacccttcc ggcccgtgt 39

<210> 5639
<211> 51
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<220>
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<223> 1 of 2 allelic variants (5640 is other entry)

<221> misc_feature
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<223> Accession number cg44014666

<400> 5639
gcgcggccct tcgggcgccc gagcccgcaa tgtcgggccc caacggagac c 51

<210> 5640
<211> 50

<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (5639 is other entry)

<221> misc_feature
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<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
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<223> Accession number cg44014666

<400> 5640
gcgcggccct tcgggcgccc gagccgcaat gtcgggcccc aacggagacc 50

<210> 5641
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5642 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44014666

<400> 5641
ttcgggcgcc cgagcccgcga atgtcgggcc ccaacggaga cctggggatg c 51

<210> 5642
<211> 50
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5641 is other entry)

<221> misc_feature
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<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
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<223> Accession number cg44014666

<400> 5642
ttcgggcgcc cgagcccgcga atgtcgggcc caacggagac ctggggatgc 50

<210> 5643
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5644 is other entry)

<221> misc_feature
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<223> Accession number cg44014666

<400> 5643
aggctctggc ctgggcactc accccctggc ttagacacct tctcaagggc t 51

<210> 5644
<211> 50
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (5643 is other entry)

<221> misc_feature
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<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44014666

<400> 5644
aggctctggc ctgggcactc acccctggct tagacacctt ctcaagggt 50

<210> 5645
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5646 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44015725

<400> 5645
aggccggctg atgcaggagg aggtgcggag ggtagagtgg tactacctcc g 51

<210> 5646
<211> 51

<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5645 is other entry)

<221> misc_feature
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<223> Accession number cg44015725

<400> 5646
aggccggctg atgcaggagg aggtgaggag ggtagagtgg tactacctcc g

51

<210> 5647
<211> 50
<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (5648 is other entry)

<221> misc_feature
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<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
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<223> Accession number cg44015725

<400> 5647
cggagcgagg aggcagcgct tttttggagg cttggcttcc cgtgcgtttc

50

<210> 5648
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (5647 is other entry)

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<223> Accession number cg44015725

<400> 5648
cggagcgagg aggcagcgct tttttggag gcttggttcc ccgtgcgttt c

51

<210> 5649
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5650 is other entry)

<221> misc_feature
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<223> Accession number cg44016438

<400> 5649
gtctatgcaa catcttccag ataatccgca acatcactga gctgggatac a 51

<210> 5650
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (5649 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44016438

<400> 5650
gtctatgcaa catcttccag ataattcgca acatcactga gctgggatac a 51

<210> 5651
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5652 is other entry)

<221> misc_feature
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<223> Accession number cg44016438

<400> 5651
catcttccag ataatccgca acatcactga gctgggatac agtcagatct t 51

<210> 5652
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (5651 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44016438

<400> 5652
catcttccag ataatccgca acatcgctga gctgggatac agtcagatct t 51

<210> 5653
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5654 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44016639

<400> 5653
ctctccacgt actgcacagg ccttggcccg ccctcaccgg ctgggccacc a 51

<210> 5654
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (5653 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44016639

<400> 5654
ctctccacgt actgcacagg ccttgtcccg ccctcaccgg ctgggccacc a 51

<210> 5655
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5656 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44017116

<400> 5655
aagtccatgat aatgatcatc actttctaaa tgagaggtct tgaggaagat g 51

<210> 5656
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (5655 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44017116

<400> 5656
aagtcctgat aatgatcatc actttttaaa tgagaggtct tgaggaagat g 51

<210> 5657
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5658 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44017116

<400> 5657
gaaaaaaaaa taattaaaaa aaaaaaccta attgtacaag ctacactgtg t 51

<210> 5658
<211> 50
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (5657 is other entry)

<221> misc_feature
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<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44017116

<400> 5658
gaaaaaaaaa taattaaaaa aaaaacctaa ttgtacaagc tacactgtgt 50

<210> 5659

<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5660 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44017354

<400> 5659
gccagagctg gtgccagggc tgaggcccaa gccagagccg agcacagaaa c 51

<210> 5660
<211> 50
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5659 is other entry)

<221> misc_feature
<222> (25)...(26)
<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44017354

<400> 5660
gccagagctg gtgccagggc tgaggccaag ccagagccga gcacagaaac 50

<210> 5661
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5662 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44018484

<400> 5661
ccattgtcaa cccctctcag gttcttgatg aagtcattcta gtttcattctt t 51

<210> 5662
<211> 51
<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (5661 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44018484

<400> 5662

ccattgtcaa cccctctcag gttctcgatg aagtcacta gtttcatctt t

51

<210> 5663

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (5664 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44018633

<400> 5663

ccaaagaagc accaaggag catctggacc accaggctgc acaccaacc t

51

<210> 5664

<211> 50

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (5663 is other entry)

<221> misc_feature

<222> (25)...(26)

<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44018633

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ccaaagaagc accaaggag catctgacca ccaggctgca caccaaccct

50

<210> 5665

<211> 51

<212> DNA

<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (5666 is other entry)

<221> misc_feature
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<223> Accession number cg44018633

<400> 5665
caaagaagca ccaagggagc atctggacca ccaggctgca caccaaccct t 51

<210> 5666
<211> 50
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<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5665 is other entry)

<221> misc_feature
<222> (25)...(26)
<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44018633

<400> 5666
caaagaagca ccaagggagc atctgaccac caggctgcac accaaccctt 50

<210> 5667
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5668 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44018633

<400> 5667
tttccagatc tttccaaagc tgatatcaat gggcagaatc caaatatcca g 51

<210> 5668
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5667 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44018633

<400> 5668
tttccagatc tttccaaagc tgataccaat gggcagaatc caaatatcca g 51

<210> 5669
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5670 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44018633

<400> 5669
ctttccaaag ctgatatcaa tgggcagaat ccaaatatcc aggtcaccat a 51

<210> 5670
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (5669 is other entry)

<221> misc_feature
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<223> Accession number cg44018633

<400> 5670
ctttccaaag ctgatatcaa tgggcggaat ccaaatatcc aggtcaccat a 51

<210> 5671
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5672 is other entry)

<221> misc_feature
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<223> Accession number cg44018634

<400> 5671
gattataaga tgcttcatcg gttcaggttt tactaccata ttagtttggt t 51

<210> 5672
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5671 is other entry)

<221> misc_feature
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<223> Accession number cg44018634

<400> 5672
gattataaga tgcttcatcg gttcatgttt tactaccata ttagtttggt t 51

<210> 5673
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5674 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44019895

<400> 5673
cttttttaaa aaaatgaggg catagtgggg gggtataaaa gtctacctag c 51

<210> 5674
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5673 is other entry)

<221> misc_feature
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<400> 5674
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<210> 5675
<211> 51

<212> DNA
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<223> 1 of 2 allelic variants (5676 is other entry)

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<223> Accession number cg44019895

<400> 5675
caaaaagcatg gtttccattg aataagatta aatatatata aaacaagtac a 51

<210> 5676
<211> 51
<212> DNA
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<223> 2 of 2 allelic variants (5675 is other entry)

<221> misc_feature
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<223> Accession number cg44019895

<400> 5676
caaaaagcatg gtttccattg aataaaatta aatatatata aaacaagtac a 51

<210> 5677
<211> 51
<212> DNA
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<223> 1 of 2 allelic variants (5678 is other entry)

<221> misc_feature
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<223> Accession number cg44020396

<400> 5677
ggcagccatg aggatccatg cccagcgtgg ggtgggacga ggcattgggag a 51

<210> 5678
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 2 of 2 allelic variants (5677 is other entry)

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<222> (0)...(0)

<223> Accession number cg44020396

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ggcagccatg aggatccatg cccagtgtgg ggtgggacga ggcattggag a

51

<210> 5679

<211> 51

<212> DNA

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<223> 1 of 2 allelic variants (5680 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44020396

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gggtgggacg aggcattggga gaatatggaa cagctttcct ggtgtcacga c

51

<210> 5680

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<223> 2 of 2 allelic variants (5679 is other entry)

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<223> Accession number cg44020396

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gggtgggacg aggcattggga gaatacggaa cagctttcct ggtgtcacga c

51

<210> 5681

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 1 of 2 allelic variants (5682 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44020396

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ctgccccagg gctcctagct gccaaagtggc agagcagggc ccggccccag a 51

<210> 5682
<211> 51
<212> DNA
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<223> 2 of 2 allelic variants (5681 is other entry)

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<223> Accession number cg44020396

<400> 5682
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<210> 5683
<211> 51
<212> DNA
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<223> 1 of 2 allelic variants (5684 is other entry)

<221> misc_feature
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agctgttctt tctcagcctt ccgccgggcc agcagctcct cctgctcacg c 51

<210> 5684
<211> 51
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<213> Homo sapiens

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<223> 2 of 2 allelic variants (5683 is other entry)

<221> misc_feature
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<223> Accession number cg44020396

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<210> 5685
<211> 51
<212> DNA

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<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (5686 is other entry)

<221> misc_feature

<222> (0)...(0)

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gcaacttcaa acaggtggag gccgagttga ttgacaagct ggacagcatg g

51

<210> 5686

<211> 51

<212> DNA

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<223> 2 of 2 allelic variants (5685 is other entry)

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51

<210> 5687

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 1 of 2 allelic variants (5688 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44020584

<400> 5687

cgagttgatt gacaagctgg acagcatggt gtcagaaggg aaaggtgacg a

51

<210> 5688

<211> 51

<212> DNA

<213> Homo sapiens

<220>

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<223> 2 of 2 allelic variants (5687 is other entry)

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<223> Accession number cg44020584

<400> 5688
cgagttgatt gacaagctgg acagcgtggt gtcagaaggg aaagtgacg a 51

<210> 5689
<211> 51
<212> DNA
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5690 is other entry)

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<223> Accession number cg44021710

<400> 5689
aagggattct tcctttgaga gaagacatgg agagcgagac cgtcgtgaca a 51

<210> 5690
<211> 51
<212> DNA
<213> Homo sapiens

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<222> (26)...(0)
<223> 2 of 2 allelic variants (5689 is other entry)

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<222> (0)...(0)
<223> Accession number cg44021710

<400> 5690
aagggattct tcctttgaga gaagatatgg agagcgagac cgtcgtgaca a 51

<210> 5691
<211> 51
<212> DNA
<213> Homo sapiens

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<222> (26)...(0)
<223> 1 of 2 allelic variants (5692 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44021758

<400> 5691

acctttctct tctgctccgg ctcaggtccc gcgggctggc tccagcagcc g

51

<210> 5692

<211> 50

<212> DNA

<213> Homo sapiens

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<223> 2 of 2 allelic variants (5691 is other entry)

<221> misc_feature

<222> (25)...(26)

<223> Nucleotide deleted between bases 25 and 26

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<222> (0)...(0)

<223> Accession number cg44021758

<400> 5692

acctttctct tctgctccgg ctcagctccg cgggctggct ccagcagccg

50

<210> 5693

<211> 51

<212> DNA

<213> Homo sapiens

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<222> (26)...(0)

<223> 1 of 2 allelic variants (5694 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44021807

<400> 5693

aggttcatcc cacatcttca catctcacc tcccaccgtc aaaacaggtc c

51

<210> 5694

<211> 51

<212> DNA

<213> Homo sapiens

<220>

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<222> (26)...(0)

<223> 2 of 2 allelic variants (5693 is other entry)

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<223> Accession number cg44021807

<400> 5694

aggttcatcc cacatcttca catcttacc tcccaccgtc aaaacaggtc c

51

<210> 5695
<211> 51
<212> DNA
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<223> 1 of 2 allelic variants (5696 is other entry)

<221> misc_feature
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<223> Accession number cg44022224

<400> 5695
gaccagcaac gccacgctt cctgggcttt gcctcgagga tccccgggga c 51

<210> 5696
<211> 50
<212> DNA
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<223> 2 of 2 allelic variants (5695 is other entry)

<221> misc_feature
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<223> Accession number cg44022224

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<210> 5697
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<223> 1 of 2 allelic variants (5698 is other entry)

<221> misc_feature
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<223> Accession number cg44023097

<400> 5697
tgccacggac ctactccgg cagtggctgc agtacacgcc gaactcgatc c 51

<210> 5698
<211> 51

<212> DNA
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<223> 2 of 2 allelic variants (5697 is other entry)

<221> misc_feature
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<400> 5698
tgccacggac ctactccgg cagtgactgc agtacacgcc gaactcgatc c 51

<210> 5699
<211> 51
<212> DNA
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<223> 1 of 2 allelic variants (5700 is other entry)

<221> misc_feature
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<223> Accession number cg44023097

<400> 5699
cactccggca gtggctgcag tacacgccga actcgatccc tttgtgaggg t 51

<210> 5700
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<212> DNA
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<223> 2 of 2 allelic variants (5699 is other entry)

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<400> 5700
cactccggca gtggctgcag tacacaccga actcgatccc tttgtgaggg t 51

<210> 5701
<211> 51
<212> DNA
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<223> 1 of 2 allelic variants (5702 is other entry)

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<223> Accession number cg44023097

<400> 5701

tccggcagtg gctgcagtag acgccgaact cgatcccttt gtgaggggtca g

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<210> 5702

<211> 51

<212> DNA

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<223> 2 of 2 allelic variants (5701 is other entry)

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<223> Accession number cg44023097

<400> 5702

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<210> 5703

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<212> DNA

<213> Homo sapiens

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<223> 1 of 2 allelic variants (5704 is other entry)

<221> misc_feature

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<400> 5703

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<212> DNA

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<223> 2 of 2 allelic variants (5703 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44023097

<400> 5704
cgatcccttt gtgaggggtca ggagggcagg agacaaactt caacacttca g 51

<210> 5705
<211> 51
<212> DNA
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<223> 1 of 2 allelic variants (5706 is other entry)

<221> misc_feature
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<223> Accession number cg44023097

<400> 5705
ctttgtgagg gtcaggagga caggagacaa acttcaacac ttcagctcgc t 51

<210> 5706
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<212> DNA
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<400> 5706
ctttgtgagg gtcaggagga caggacacaa acttcaacac ttcagctcgc t 51

<210> 5707
<211> 51
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<223> 1 of 2 allelic variants (5708 is other entry)

<221> misc_feature
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<223> Accession number cg44023097

<400> 5707
tcaggaggac aggagacaaa cttcaacact tcagctcgc tctctctcag a 51

<210> 5708
<211> 51
<212> DNA

<213> Homo sapiens

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<222> (26)...(0)

<223> 2 of 2 allelic variants (5707 is other entry)

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<223> Accession number cg44023097

<400> 5708

tcaggaggac aggagacaaa cttcagcact tcagctcgct tctctctcag a

51

<210> 5709

<211> 51

<212> DNA

<213> Homo sapiens

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<222> (26)...(0)

<223> 1 of 2 allelic variants (5710 is other entry)

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51

<210> 5710

<211> 51

<212> DNA

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51

<210> 5711

<211> 51

<212> DNA

<213> Homo sapiens

<220>

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<222> (26)...(0)

<223> 1 of 2 allelic variants (5712 is other entry)

<221> misc_feature
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<223> Accession number cg44023776

<400> 5711
tacacaaaat gatcattcca taaatattta catgacaagg gaaaaaatgg a 51

<210> 5712
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<212> DNA
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<220>
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<223> 2 of 2 allelic variants (5711 is other entry)

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tacacaaaat gatcattcca taaatcttta catgacaagg gaaaaaatgg a 51

<210> 5713
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<223> 1 of 2 allelic variants (5714 is other entry)

<221> misc_feature
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<223> Accession number cg44023776

<400> 5713
ggtgtgataa tcctttctca tgacacttta gtttaggaatc atgcaagctt t 51

<210> 5714
<211> 51
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<223> 2 of 2 allelic variants (5713 is other entry)

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<400> 5714

gggtgtgataa tccttttctca tgacatttta gtttaggaatc atgcaagctt t

51

<210> 5715

<211> 51

<212> DNA

<213> Homo sapiens

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<222> (26)...(0)

<223> 1 of 2 allelic variants (5716 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44023776

<400> 5715

taaaatgaaa ataattgtag aagcgtaac taaaatattc cattttagtt t

51

<210> 5716

<211> 50

<212> DNA

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<221> misc_feature

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<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44023776

<400> 5716

taaaatgaaa ataattgtag aagcgtaact aaaatattcc attttagttt

50

<210> 5717

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

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<223> Accession number cg44023776

<400> 5717

aaaatgaaaa taattgtaga agcgtaact aaaatattcc attttagttt t

51

<210> 5718
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<221> misc_feature
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<221> misc_feature
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<400> 5718
aaaatgaaaa taattgtaga agcgtaacta aaatattcca ttttagtttt

50

<210> 5719
<211> 51
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<221> misc_feature
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<223> Accession number cg44023776

<400> 5719
gtttaatttt cagtatcaga tcttttcctt catctctcat gatgaaacta a

51

<210> 5720
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gtttaatttt cagtatcaga tcttttcctt catctctcat gatgaaacta a

51

<210> 5721
<211> 51

<212> DNA
<213> Homo sapiens

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<400> 5721
cagttgtcca tgctcaaggc atctgccttc cttcggagtc gatcatcacg g 51

<210> 5722
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<223> 2 of 2 allelic variants (5721 is other entry)

<221> misc_feature
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<400> 5722
cagttgtcca tgctcaaggc atctgacttc cttcggagtc gatcatcacg g 51

<210> 5723
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5724 is other entry)

<221> misc_feature
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<223> Accession number cg44023776

<400> 5723
atctgccttc cttcggagtc gatcatcacg gtatactttt gctgcatact g 51

<210> 5724
<211> 51
<212> DNA
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<220>
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<223> 2 of 2 allelic variants (5723 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44023776

<400> 5724

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51

<210> 5725

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 1 of 2 allelic variants (5726 is other entry)

<221> misc_feature

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<223> Accession number cg44023776

<400> 5725

gaagtccttg ttgacataat aaacaacacc caagttttct gtttgcattt t

51

<210> 5726

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 2 of 2 allelic variants (5725 is other entry)

<221> misc_feature

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<223> Accession number cg44023776

<400> 5726

gaagtccttg ttgacataat aaacaccacc caagttttct gtttgcattt t

51

<210> 5727

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 1 of 2 allelic variants (5728 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44023886

<400> 5727
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<210> 5728
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<213> Homo sapiens

<220>
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<223> Accession number cg44023886

<400> 5728
ttcagtgtaa acttgacacg ttccgctgcg agcagccacg ggtctggta c 50

<210> 5729
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<223> 1 of 2 allelic variants (5730 is other entry)

<221> misc_feature
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<223> Accession number cg44024241

<400> 5729
gggggactat gaaatccacg atgggatgaa cctggagctt tattatcaat a 51

<210> 5730
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 2 of 2 allelic variants (5729 is other entry)

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<223> Accession number cg44024241

<400> 5730
gggggactat gaaatccacg atggggtgaa cctggagctt tattatcaat a 51

<210> 5731
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 1 of 2 allelic variants (5732 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44024241

<400> 5731
cccgtttcc tctcccatcc tcatcccca cactgggata gatgcttggt t 51

<210> 5732
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 2 of 2 allelic variants (5731 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44024241

<400> 5732
cccgtttcc tctcccatcc tcatctccca cactgggata gatgcttggt t 51

<210> 5733
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (5734 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44024241

<400> 5733
ccccacact gggatagatg cttgtttgta aaaactcacc ttaataaaga c 51

<210> 5734
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5733 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44024241

<400> 5734
ccccacact gggatagatg cttgtctgta aaaactcacc ttaataaaga c 51

<210> 5735
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5736 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44024675

<400> 5735
gtcactggaa ttattattat gagaatgtct ttgggccccca tttgtgccaa a 51

<210> 5736
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5735 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44024675

<400> 5736
gtcactggaa ttattattat gagaacgtct ttgggccccca tttgtgccaa a 51

<210> 5737
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
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<223> 1 of 2 allelic variants (5738 is other entry)

<221> misc_feature

<222> (0)...(0)
<223> Accession number cg44025359

<400> 5737
ggttagagga gacagaaaac aaagcccagc tcttccaagc tcaccaccta g 51

<210> 5738
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5737 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44025359

<400> 5738
ggttagagga gacagaaaac aaagcacagc tcttccaagc tcaccaccta g 51

<210> 5739
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5740 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44025363

<400> 5739
gtcaagttca tattgtatca cattgttggt ggatgacact ggctccagat a 51

<210> 5740
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (5739 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44025363

<400> 5740
gtcaagttca tattgtatca cattgctggt ggatgacact ggctccagat a 51

<210> 5741
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5742 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44025788

<400> 5741
ctaataaggaa aacgaagacc caagatTTTT ttttaaaatt aaggttatTT t 51

<210> 5742
<211> 50
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5741 is other entry)

<221> misc_feature
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<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44025788

<400> 5742
ctaataaggaa aacgaagacc caagatTTTT ttttaaaatta aggttatTTT 50

<210> 5743
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5744 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44026589

<400> 5743
tgacactgac tttctaaata ttccagaact atttccctga acgtgaaggt c 51

<210> 5744
<211> 51

<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5743 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44026589

<400> 5744
tgacactgac tttctaaata ttccacaact atttcctga acgtgaaggt c 51

<210> 5745
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5746 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44026882

<400> 5745
ttttgagatt cttctcaata atttcggcaa atctggttg cgatcatctca t 51

<210> 5746
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (5745 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44026882

<400> 5746
ttttgagatt cttctcaata atttcagcaa atctggttg cgatcatctca t 51

<210> 5747
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)

<223> 1 of 2 allelic variants (5748 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44027139

<400> 5747

gtgcagcggc tcacacctgt aatcctagca ctttaggaga ctgaggtggg a

51

<210> 5748

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (5747 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44027139

<400> 5748

gtgcagcggc tcacacctgt aatcccagca ctttaggaga ctgaggtggg a

51

<210> 5749

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (5750 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44027365

<400> 5749

agaagtgtccc tatctgcaga caggcgatca cccgggtgat acccctgtac a

51

<210> 5750

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (5749 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44027365

<400> 5750
agaagtgtccc tatctgcaga caggcaatca cccgggtgat acccctgtac a 51

<210> 5751
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5752 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44027572

<400> 5751
ctggattaag ggtcagggcc ggtggctgtg ggaggtgaca ctgagccgct c 51

<210> 5752
<211> 50
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5751 is other entry)

<221> misc_feature
<222> (25)...(26)
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<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44027572

<400> 5752
ctggattaag ggtcagggcc ggtgggtgtgg gaggtgacac tgagccgctc 50

<210> 5753
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5754 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44027650

<400> 5753
ttacaagaga tttcgactgt ggaagtaaca aatactttta agaaaacaga t 51

<210> 5754
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5753 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44027650

<400> 5754
ttacaagaga ttctgactgt ggaagcaaca aatactttta agaaaacaga t 51

<210> 5755
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5756 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44029982

<400> 5755
ggggtgcaca tcatctggtg gtgatggtgg ctgtagggga tgttggtctc c 51

<210> 5756
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5755 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44029982

<400> 5756
ggggtgcaca tcatctggtg gtgatcgtgg ctgtagggga tgttggtctc c 51

<210> 5757
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5758 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44030068

<400> 5757
agaaatattc attgaagttg attatagagg aaagaggaaa gttggcatat t 51

<210> 5758
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5757 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44030068

<400> 5758
agaaatattc attgaagttg attatggagg aaagaggaaa gttggcatat t 51

<210> 5759
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5760 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44030287

<400> 5759
ttgccttttg aagaagagcc cgcattatga gtggacggca gacagctata t 51

<210> 5760
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5759 is other entry)

<221> misc_feature

<222> (0)...(0)
<223> Accession number cg44030287

<400> 5760
ttgccttttg aagaagagcc cgcataatga gtggacggca gacagctata t 51

<210> 5761
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5762 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44030433

<400> 5761
gatcaaagtc tattttgcat aaaatgtcca ataattaaat attgttataa a 51

<210> 5762
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (5761 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44030433

<400> 5762
gatcaaagtc tattttgcat aaaatctcca ataattaaat attgttataa a 51

<210> 5763
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5764 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44030437

<400> 5763
cagctgccca ccaagactca agcaacgtaa gagtcctctc cccagactgg c 51

<210> 5764
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (5763 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44030437

<400> 5764
cagcctgccca ccaagactca agcaatgtaa gagtcatctc cccagactgg c 51

<210> 5765
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5766 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44030903

<400> 5765
ggcggttcggg aagggggccgc ggccggtccg gggccgtcgc gcttgggggc c 51

<210> 5766
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5765 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44030903

<400> 5766
ggcggttcggg aagggggccgc ggccgatccg gggccgtcgc gcttgggggc c 51

<210> 5767
<211> 51
<212> DNA
<213> Homo sapiens

<220>

<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5768 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44031455

<400> 5767
cctgctcccc aactacagca gggctcttggg tcctgggtct gagggtttat t 51

<210> 5768
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (5767 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44031455

<400> 5768
cctgctcccc aactacagca gggctcctggg tcctgggtct gagggtttat t 51

<210> 5769
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5770 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44031677

<400> 5769
ccagtggcat tccagacttc agatcgccgg ggagtggcct gtacagcaac c 51

<210> 5770
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (5769 is other entry)

<221> misc_feature
<222> (0)...(0)

<223> Accession number cg44031677

<400> 5770

ccagtggcat tccagacttc agatctccgg ggagtggcct gtacagcaac c

51

<210> 5771

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (5772 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44032154

<400> 5771

ggcctctctc tgcctacgga ggggggggaa tctaaccct ctgccctggc t

51

<210> 5772

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (5771 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44032154

<400> 5772

ggcctctctc tgcctacgga ggggggtggaa tctaaccct ctgccctggc t

51

<210> 5773

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (5774 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44032208

<400> 5773

catttttatg tactcatctg ctgtgaaaag tcttttaggtt cattaaaaa a

51

<210> 5774

<211> 50
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (5773 is other entry)

<221> misc_feature
<222> (25)...(26)
<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44032208

<400> 5774
catttttatg tactcatctg ctgtgaaagt ctttaggttc attaaaaaaaa

50

<210> 5775
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5776 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44032208

<400> 5775
tttttatgta ctcatctgct gtgaaaagtc ttttaggttca ttaaaaaaac a

51

<210> 5776
<211> 50
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (5775 is other entry)

<221> misc_feature
<222> (25)...(26)
<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44032208

<400> 5776
tttttatgta ctcatctgct gtgaaagtct ttaggttcat taaaaaaca

50

<210> 5777
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5778 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44032208

<400> 5777
ttttatgtac tcattctgctg tgaaaagtct ttaggttcatt taaaaaaaca g 51

<210> 5778
<211> 50
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5777 is other entry)

<221> misc_feature
<222> (25)...(26)
<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44032208

<400> 5778
ttttatgtac tcattctgctg tgaaagtctt taggttcatt aaaaaaacag 50

<210> 5779
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5780 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44032208

<400> 5779
tagaaatgat cttagatcta atatagtgat tttaagcatc ccgtcaaagg c 51

<210> 5780

<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (5779 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44032208

<400> 5780
tagaaatgat cttagatcta atatactgat ttttaagcatc ccgtcaaagg c

51

<210> 5781
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5782 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44032216

<400> 5781
ctgagtattt tatgctatat gtgtgtgagt atatatatgt gtgtatatct a

51

<210> 5782
<211> 50
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (5781 is other entry)

<221> misc_feature
<222> (25)...(26)
<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44032216

<400> 5782
ctgagtattt tatgctatat gtgtggagta tatatatgtg tgtatatcta

50

<210> 5783
<211> 51
<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (5784 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44032244

<400> 5783

cttgagcctg ggaggtgaag gttgcagtga gccaaagatca cgccactgca c

51

<210> 5784

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (5783 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44032244

<400> 5784

cttgagcctg ggaggtgaag gttgcggtga gccaaagatca cgccactgca c

51

<210> 5785

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (5786 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44034193

<400> 5785

gggccatctc agtgggctgg tacacactgg tcagcagctg cccattgtac c

51

<210> 5786

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (5785 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44034193

<400> 5786
gggccatctc agtgggctgg tacacgctgg tcagcagctg cccattgtac c 51

<210> 5787
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5788 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44034193

<400> 5787
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<210> 5788
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5787 is other entry)

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<223> Accession number cg44034193

<400> 5788
gctggtacac actggtcagc agctgacat tgtaccgct gtatggtgac a 51

<210> 5789
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (5790 is other entry)

<221> misc_feature
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<223> Accession number cg44034193

<400> 5789

cactggtcag cagctgccca ttgtaccgcg tgtatgggtga caccggcctc t

51

<210> 5790

<211> 51

<212> DNA

<213> Homo sapiens

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<222> (26)...(0)

<223> 2 of 2 allelic variants (5789 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44034193

<400> 5790

cactggtcag cagctgccca ttgtaccgcg tgtatgggtga caccggcctc t

51

<210> 5791

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 1 of 2 allelic variants (5792 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44034691

<400> 5791

gagtggccac ctgcgaagac agggccgtca ttatgggcca gagggctgct c

51

<210> 5792

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (5791 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44034691

<400> 5792

gagtggccac ctgcgaagac agggctgtca ttatgggcca gagggctgct c

51

<210> 5793

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 1 of 2 allelic variants (5794 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44034774

<400> 5793
gtggtactcc aggtcctctg tcagccgctg gagcatggac aggggctcat t 51

<210> 5794
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (5793 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44034774

<400> 5794
gtggtactcc aggtcctctg tcagctgctg gagcatggac aggggctcat t 51

<210> 5795
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 1 of 2 allelic variants (5796 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44034830

<400> 5795
ggggaccag gcattccggt cccccctggc ggtgagggtta ccaatggcct c 51

<210> 5796
<211> 50
<212> DNA
<213> Homo sapiens

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<221> misc_feature
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<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44034830

<400> 5796
ggggacccag gcattccggt ccccttggcg gtgaggggtac caatggcctc 50

<210> 5797
<211> 51
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<213> Homo sapiens

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<223> 1 of 2 allelic variants (5798 is other entry)

<221> misc_feature
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<223> Accession number cg44034830

<400> 5797
atggccttgg cagggacatc acagcgaaac acatggcact tgagcataca g 51

<210> 5798
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 2 of 2 allelic variants (5797 is other entry)

<221> misc_feature
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<223> Accession number cg44034830

<400> 5798
atggccttgg cagggacatc acagcaaaac acatggcact tgagcataca g 51

<210> 5799
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (5800 is other entry)

<221> misc_feature
<222> (0)...(0)

<223> Accession number cg44034830

<400> 5799

gtagaagctg gttagaggca ggagcctgca ggaggctgga aagtcaggct a

51

<210> 5800

<211> 50

<212> DNA

<213> Homo sapiens

<220>

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<222> (26)...(0)

<223> 2 of 2 allelic variants (5799 is other entry)

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<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44034830

<400> 5800

gtagaagctg gttagaggca ggagctgcag gaggtggaa agtcaggcta

50

<210> 5801

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (5802 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44034889

<400> 5801

ggccgtccgg gactgcgtgg agaggacgg gctccacagc gtggtggatg a

51

<210> 5802

<211> 51

<212> DNA

<213> Homo sapiens

<220>

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<222> (26)...(0)

<223> 2 of 2 allelic variants (5801 is other entry)

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<222> (0)...(0)

<223> Accession number cg44034889

<400> 5802
ggcgcgtccgg gactgcgtgg agaggaacgg gctccacagc gtggtggatg a 51

<210> 5803
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5804 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44034889

<400> 5803
cgccggccgc tgcgctgccc gggacggaga cagaggggcc ggacctccca g 51

<210> 5804
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5803 is other entry)

<221> misc_feature
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<223> Accession number cg44034889

<400> 5804
cgccggccgc tgcgctgccc gggacagaga cagaggggcc ggacctccca g 51

<210> 5805
<211> 50
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5806 is other entry)

<221> misc_feature
<222> (25)...(26)
<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44035239

<400> 5805
ttttatcagc tatatatata tatatgagaa tatatatata ttttgttgtt 50

<210> 5806
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5805 is other entry)

<221> misc_feature
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<223> Accession number cg44035239

<400> 5806
ttttatcagc tatatatata tatatagaga atatatatat attttgtgt t 51

<210> 5807
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5808 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44035419

<400> 5807
actccaagac cgtgagtccc ctagaagtta ctcatccact ttgactgaca t 51

<210> 5808
<211> 51
<212> DNA
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<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5807 is other entry)

<221> misc_feature
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<223> Accession number cg44035419

<400> 5808
actccaagac cgtgagtccc ctagagggtta ctcatccact ttgactgaca t 51

<210> 5809
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5810 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44035528

<400> 5809
ggagctcggg gtgggggtcgt gggacggtgc gcgatagcat cagtccacag g 51

<210> 5810
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5809 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44035528

<400> 5810
ggagctcggg gtgggggtcgt gggaccgtgc gcgatagcat cagtccacag g 51

<210> 5811
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5812 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44036050

<400> 5811
agctcgaagg agccaaagcc agcggctgcg aaactttctt cgtatcttcc c 51

<210> 5812
<211> 50
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (5811 is other entry)

<221> misc_feature

<222> (25)...(26)
<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
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<223> Accession number cg44036050

<400> 5812
agctcgaagg agccaaagcc agcgggtgcga aactttcttc gtatcttccc 50

<210> 5813
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5814 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44036050

<400> 5813
ggtcgatgta gaccttagta ccatgtccga tgagatactg tccgtgtttg t 51

<210> 5814
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5813 is other entry)

<221> misc_feature
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<223> Accession number cg44036050

<400> 5814
ggtcgatgta gaccttagta ccatgcccga tgagatactg tccgtgtttg t 51

<210> 5815
<211> 50
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5816 is other entry)

<221> misc_feature
<222> (25)...(26)
<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44036050

<400> 5815
taagaccaaa aacaaaactc aaaaaccttc aatatgaagg cagcagctgg 50

<210> 5816
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5815 is other entry)

<221> misc_feature
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<223> Accession number cg44036050

<400> 5816
taagaccaaa aacaaaactc aaaaaacctt caatatgaag gcagcagctg g 51

<210> 5817
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5818 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44036247

<400> 5817
gtacaaaaat ttacataaca agaggaaaaa taggcagtgc agcaccttta g 51

<210> 5818
<211> 50
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5817 is other entry)

<221> misc_feature
<222> (25)...(26)
<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature

<222> (0)...(0)
<223> Accession number cg44036247

<400> 5818
gtacaaaaat ttacataaca agaggaaaat aggcagtgcg gcacctttag 50

<210> 5819
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5820 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44126480

<400> 5819
gtggtgcatc tatctatggc aaacaatttg aagatgaact tcacccagac t 51

<210> 5820
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5819 is other entry)

<221> misc_feature
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<223> Accession number cg44126480

<400> 5820
gtggtgcatc tatctatggc aaacagtttg aagatgaact tcacccagac t 51

<210> 5821
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5822 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44126938

<400> 5821
ggtacttttt cgcccgctct cgatgaaatc cctgatggc aagaactgtt t 51

<210> 5822
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5821 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44126938

<400> 5822
gggtacttttt cgcccgctcct cgatggaatc cctgatggc aagaactggt t 51

<210> 5823
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5824 is other entry)

<221> misc_feature
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<223> Accession number cg44126938

<400> 5823
gggtcccgcga acggcggata cttcgtcggt gacacgcttc gtgcgtgggt g 51

<210> 5824
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5823 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44126938

<400> 5824
gggtcccgcga acggcggata cttcgccggt gacacgcttc gtgcgtgggt g 51

<210> 5825
<211> 51
<212> DNA
<213> Homo sapiens

<220>

<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5826 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44126946

<400> 5825
ccacgttctt ggtcacgctg gatccccaaa cgcataccgg ggtagcagcc c 51

<210> 5826
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5825 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44126946

<400> 5826
ccacgttctt ggtcacgctg gatcctcaaa cgcataccgg ggtagcagcc c 51

<210> 5827
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5828 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44127370

<400> 5827
ggcgcccaa gacaccgtgc tcgtcacagg tgtgggacaa caccaggtgc g 51

<210> 5828
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
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<223> 2 of 2 allelic variants (5827 is other entry)

<221> misc_feature
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<223> Accession number cg44127370

<400> 5828

ggcgcccaaa gacaccgtgc tcgtcgcagg tgtgggacaa caccagggtgc g

51

<210> 5829

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (5830 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44127374

<400> 5829

gttgggctcc tgcggcgttt gttgatcggg tgctcggacc cttacagccc a

51

<210> 5830

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (5829 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44127374

<400> 5830

gttgggctcc tgcggcgttt gttgagcggg tgctcggacc cttacagccc a

51

<210> 5831

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (5832 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44127374

<400> 5831

ctgatcaggc cccggccttg ctgtgttcgc tggagcgacg gatcgagtgg c

51

<210> 5832

<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5831 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44127374

<400> 5832
ctgatcaggc cccggccttg ctgtgctcgc tggagcgcgc gatcgagtgg c 51

<210> 5833
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5834 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44127378

<400> 5833
gtgcgccatc gtcacgaagc agctggcccc tcagccaggg aaaagcatcc g 51

<210> 5834
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (5833 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44127378

<400> 5834
gtgcgccatc gtcacgaagc agctgtcccc tcagccaggg aaaagcatcc g 51

<210> 5835
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature

<222> (26)...(0)
<223> 1 of 2 allelic variants (5836 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44127419

<400> 5835
tcccctcagc aatggcctca gaaacctcgg cacgaatctc ctcgatacgc c 51

<210> 5836
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (5835 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44127419

<400> 5836
tcccctcagc aatggcctca gaaacctcgg cacgaatctc ctcgatacgc c 51

<210> 5837
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5838 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44127419

<400> 5837
gacgcaagcc ctcgccaccc tcttcaacaa caaagaggcc cctcaccacg t 51

<210> 5838
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (5837 is other entry)

<221> misc_feature
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<223> Accession number cg44127419

<400> 5838
gacgcaagcc ctcgccaccc tcttcgacaa caaagaggcc cctcaccacg t 51

<210> 5839
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5840 is other entry)

<221> misc_feature
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<223> Accession number cg44127419

<400> 5839
gaacgtcacc agcctcgacg acgagaccgg cccgggtacg ggtcttctcc c 51

<210> 5840
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (5839 is other entry)

<221> misc_feature
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<223> Accession number cg44127419

<400> 5840
gaacgtcacc agcctcgacg acgagtccgg cccgggtacg ggtcttctcc c 51

<210> 5841
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5842 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44127465

<400> 5841
ggccctctg ggccgcaggg accccctggg gacagccgcc tcctgtccac g 51

<210> 5842
<211> 50

<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (5841 is other entry)

<221> misc_feature
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<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44127465

<400> 5842
ggccctcctg ggccgcaggg acccctgggg acagccgcct cctgtccacg 50

<210> 5843
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5844 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44127465

<400> 5843
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<210> 5844
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (5843 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44127465

<400> 5844
gtgacagggg cctgtctggg ccaccgggtc atcctggggc acctggccct c 51

<210> 5845
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5846 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44127475

<400> 5845
caccactgta tacaccgacg atggtaggaa aacgtggcaa cggggaatcc c 51

<210> 5846
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (5845 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44127475

<400> 5846
caccactgta tacaccgacg atggtgggaa aacgtggcaa cggggaatcc c 51

<210> 5847
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5848 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44127475

<400> 5847
cactgtatac accgacgatg gtaggaaaac gtggcaacgg ggaatcccga t 51

<210> 5848
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 2 of 2 allelic variants (5847 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44127475

<400> 5848
cactgtatac accgacgatg gtagggaaac gtggcaacgg ggaatcccga t 51

<210> 5849
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (5850 is other entry)

<221> misc_feature
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<223> Accession number cg44127475

<400> 5849
gggaacagct tcctgcgctc attgtcgtct cagcagatga ccgcaggcgt c 51

<210> 5850
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5849 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44127475

<400> 5850
gggaacagct tcctgcgctc attgttgtct cagcagatga ccgcaggcgt c 51

<210> 5851
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5852 is other entry)

<221> misc_feature
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<223> Accession number cg44127475

<400> 5851
cgctgacatg gcattgccca tgtggcagtt atccatcgat cctcgaggac g 51

<210> 5852
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (5851 is other entry)

<221> misc_feature
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<223> Accession number cg44127475

<400> 5852
cgtcgacatg gcattgccca tgtggtagtt atccatcgat cctcgaggac g 51

<210> 5853
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5854 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44127493

<400> 5853
catctacgac cgccgagacc tgacgctcgt tggccgccga gatgacaacg a 51

<210> 5854
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5853 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44127493

<400> 5854
catctacgac cgccgagacc tgacgttcgt tggccgccga gatgacaacg a 51

<210> 5855
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (5856 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44127493

<400> 5855
aagcccttct tgccaagggc agcgtgccc gctacgcggg caagctcgat a 51

<210> 5856
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (5855 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44127493

<400> 5856
aagcccttct tgccaagggc agcgtacgcc gctacgcggg caagctcgat a 51

<210> 5857
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
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<223> 1 of 2 allelic variants (5858 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44127493

<400> 5857
cttcttgcca agggcagcgt gcgccgtac gcgggcaagc tcgatagcat a 51

<210> 5858
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5857 is other entry)

<221> misc_feature

<222> (0)...(0)
<223> Accession number cg44127493

<400> 5858
cttcttgcca agggcagcgt gcgccactac gcgggcaagc tcgatagcat a 51

<210> 5859
<211> 35
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (10)...(0)
<223> 1 of 2 allelic variants (5860 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44127499

<400> 5859
tccggacttt ggtctggcct ctctgacgg gccag 35

<210> 5860
<211> 35
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (10)...(0)
<223> 2 of 2 allelic variants (5859 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44127499

<400> 5860
tccggacttc ggtctggcct ctctgacgg gccag 35

<210> 5861
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5862 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44127556

<400> 5861
aggccgcctc aaggcggagt gcggtttacc tccggccgac cccgcccgta a 51

<210> 5862
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (5861 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44127556

<400> 5862
aggccgcctc aaggcggagt gcggtctacc tccggccgac cccgcccgtg a 51

<210> 5863
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5864 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44127556

<400> 5863
ccgacccgc ccgtgaggct gagcagatcg cgcggttgcg gcagttagcg g 51

<210> 5864
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5863 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44127556

<400> 5864
ccgacccgc ccgtgaggct gagcaaatcg cgcggttgcg gcagttagcg g 51

<210> 5865
<211> 51
<212> DNA
<213> Homo sapiens

<220>

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<222> (26)...(0)
<223> 1 of 2 allelic variants (5866 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44127556

<400> 5865
tggccgaggt ggtgcgtcac cacgaagcta ttgctgacga ttctggcgac g 51

<210> 5866
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (5865 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44127556

<400> 5866
tggccgaggt ggtgcgtcac cacgaggcta ttgctgacga ttctggcgac g 51

<210> 5867
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5868 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44127556

<400> 5867
gtcaccacga agctattgct gacgattctg gcgacgactc tggagtggcg g 51

<210> 5868
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (5867 is other entry)

<221> misc_feature
<222> (0)...(0)

<223> Accession number cg44127556

<400> 5868

gtcaccacga agctattgct gacgagtctg gcgacgactc tggagtggcg g

51

<210> 5869

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (5870 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44127556

<400> 5869

tgctgacgat tctggcgacg actctggagt ggcggatacg ggggagggcg a

51

<210> 5870

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (5869 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44127556

<400> 5870

tgctgacgat tctggcgacg actctagagt ggcggatacg ggggagggcg a

51

<210> 5871

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (5872 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44127560

<400> 5871

gcaaagtcgt ttctggcatg aatgatgctc agatgcgggc gctgcgtcgc g

51

<210> 5872

<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (5871 is other entry)

<221> misc_feature
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<223> Accession number cg44127560

<400> 5872
gcaaagtcgt ttctggcatg aatgacgctc agatgcgggc gctgcgtcgc g 51

<210> 5873
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5874 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44127560

<400> 5873
cgtttctggc atgaatgatg ctcatgatgcg ggcgctgcgt cgcgaccacg t 51

<210> 5874
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5873 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44127560

<400> 5874
cgtttctggc atgaatgatg ctcatgtgcg ggcgctgcgt cgcgaccacg t 51

<210> 5875
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5876 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44127560

<400> 5875
gggcgctgcg tcgcgaccac gtctcaatgg tttccagca cttcgacta c 51

<210> 5876
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5875 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44127560

<400> 5876
gggcgctgcg tcgcgaccac gtctcgatgg tttccagca cttcgacta c 51

<210> 5877
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5878 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44127560

<400> 5877
gtcgcgacca cgtctcaatg gtttccagc acttcgcact acttcctcat c 51

<210> 5878
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (5877 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44127560

<400> 5878
gtcgcgacca cgtctcaatg gtttttcagc acttcgcact acttcctcat c 51

<210> 5879
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5880 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44127560

<400> 5879
agtccaccct catcaggatg attaacgggc tgtgggtcccc gtctgagggt a 51

<210> 5880
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5879 is other entry)

<221> misc_feature
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<223> Accession number cg44127560

<400> 5880
agtccaccct catcaggatg attaatgggc tgtgggtcccc gtctgagggt a 51

<210> 5881
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5882 is other entry)

<221> misc_feature
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<223> Accession number cg44127560

<400> 5881
agggtaccgt tgaggttgcc ggcaaagtcg tttctggcat gaatgatgct c 51

<210> 5882
<211> 51

<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5881 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44127560

<400> 5882
agggtaccgt tgagggtgcc ggcaaggctcg ttcttgcat gaatgatgct c 51

<210> 5883
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5884 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44127560

<400> 5883
gagggtgccg gcaaagtcgt ttctggcatg aatgatgctc agatgcgggc g 51

<210> 5884
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (5883 is other entry)

<221> misc_feature
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<223> Accession number cg44127560

<400> 5884
gagggtgccg gcaaagtcgt ttctgacatg aatgatgctc agatgcgggc g 51

<210> 5885
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)

<223> 1 of 2 allelic variants (5886 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44127562

<400> 5885

ggtcatcctt gcttgggagc tgagccaggg cccgcacgtc atcccgattc c

51

<210> 5886

<211> 51

<212> DNA

<213> Homo sapiens

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<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (5885 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44127562

<400> 5886

ggtcatcctt gcttgggagc tgagcaaggg cccgcacgtc atcccgattc c

51

<210> 5887

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (5888 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44127562

<400> 5887

ttgggagctg agccaggggc cgcacgtcat cccgattccc gggctctcacc g

51

<210> 5888

<211> 51

<212> DNA

<213> Homo sapiens

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<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (5887 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44127562

<400> 5888
ttgggagctg agccagggcc cgcacatcat cccgattccc gggcttcacc g 51

<210> 5889
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5890 is other entry)

<221> misc_feature
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<223> Accession number cg44127562

<400> 5889
cgcacgtcat cccgattccc gggcttcacc gttcccagac gattcttgac t 51

<210> 5890
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5889 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44127562

<400> 5890
cgcacgtcat cccgattccc ggggtcccacc gttcccagac gattcttgac t 51

<210> 5891
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5892 is other entry)

<221> misc_feature
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<223> Accession number cg44127562

<400> 5891
ctcacggttc ccagacgatt cttgactccc tcagggtcgt cgacgtcacg c 51

<210> 5892
<211> 51
<212> DNA

<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5891 is other entry)

<221> misc_feature
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<400> 5892
ctcacggttc ccagacgatt cttgattccc tcagggtcgt cgacgtcacg c 51

<210> 5893
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5894 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44127562

<400> 5893
cctcaggtcc gtcgacgtca cgctggacga cgaggagctt gcccagcttc c 51

<210> 5894
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5893 is other entry)

<221> misc_feature
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<223> Accession number cg44127562

<400> 5894
cctcaggtcc gtcgacgtca cgctgaacga cgaggagctt gcccagcttc c 51

<210> 5895
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5896 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44127564

<400> 5895
gttgcatgc cgtaggcgcg ggcggtttgg tcgtgcaact caatgacctg g 51

<210> 5896
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5895 is other entry)

<221> misc_feature
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<400> 5896
gttgcatgc cgtaggcgcg ggcgggttgg tcgtgcaact caatgacctg g 51

<210> 5897
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5898 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44127564

<400> 5897
agggtgtcca atagccacgt ggaccaatgg acgtcctttt ggtaacctg c 51

<210> 5898
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5897 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44127564

<400> 5898

agggtgtcca atagccacgt ggaccgatgg acgtcctttt ggtaacctgg c 51

<210> 5899
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5900 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44127564

<400> 5899
gtccaatagc cacgtggacc aatggacgtc cttttggtaa cctggcttgc t 51

<210> 5900
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (5899 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44127564

<400> 5900
gtccaatagc cacgtggacc aatggcgctc cttttggtaa cctggcttgc t 51

<210> 5901
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<221> misc_feature
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accaatggac gtccttttgg taacctggct tgctcggacg gcatagccgc c 51

<210> 5902
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 2 of 2 allelic variants (5901 is other entry)

<221> misc_feature
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<223> Accession number cg44127564

<400> 5902
accaatggac gtccttttgg taaccgggct tgctcggacg gcatagccgc c 51

<210> 5903
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 1 of 2 allelic variants (5904 is other entry)

<221> misc_feature
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<223> Accession number cg44127564

<400> 5903
cccacgtgaa atagccaatg cggcagcccc aggtttgaag gtgtccattg c 51

<210> 5904
<211> 50
<212> DNA
<213> Homo sapiens

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<221> misc_feature
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<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
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<223> Accession number cg44127564

<400> 5904
cccacgtgaa atagccaatg cggcacccca gggtttgaagg tgtccattgc 50

<210> 5905
<211> 51
<212> DNA
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<223> 1 of 2 allelic variants (5906 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44127564

<400> 5905
ccccagggttt gaaggtgtcc attgcgccgg tgcgagaccg ggtgccctcc g 51

<210> 5906
<211> 51
<212> DNA
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<223> 2 of 2 allelic variants (5905 is other entry)

<221> misc_feature
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<223> Accession number cg44127564

<400> 5906
ccccagggttt gaaggtgtcc attgcaccgg tgcgagaccg ggtgccctcc g 51

<210> 5907
<211> 51
<212> DNA
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<223> 1 of 2 allelic variants (5908 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44127568

<400> 5907
acgatgtgga tcgtgtctct tggagttttc atctttgtcc agcagatggt c 51

<210> 5908
<211> 50
<212> DNA
<213> Homo sapiens

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<223> 2 of 2 allelic variants (5907 is other entry)

<221> misc_feature
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<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44127568

<400> 5908

acgatgtgga tcgtgctcct tggagtttca tctttgtcca gcagatgttc

50

<210> 5909

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 1 of 2 allelic variants (5910 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44127585

<400> 5909

tccagctgga gcggatggat gagagtgatg acggtgacgc cactaccggc a

51

<210> 5910

<211> 51

<212> DNA

<213> Homo sapiens

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<222> (26)...(0)

<223> 2 of 2 allelic variants (5909 is other entry)

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<222> (0)...(0)

<223> Accession number cg44127585

<400> 5910

tccagctgga gcggatggat gagagagatg acggtgacgc cactaccggc a

51

<210> 5911

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (5912 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44127585

<400> 5911
caaggacggc ctttgacccc cgcacctcgc actcggcgat ggacaaaatg t 51

<210> 5912
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 2 of 2 allelic variants (5911 is other entry)

<221> misc_feature
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<223> Accession number cg44127585

<400> 5912
caaggacggc ctttgacccc cgcacctcgc actcggcgat ggacaaaatg t 51

<210> 5913
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (5914 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44127585

<400> 5913
cctttgaccc cccgacctcg cactcggcga tggacaaaat gtggcgtaat g 51

<210> 5914
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5913 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44127585

<400> 5914
cctttgaccc cccgacctcg cactcagcga tggacaaaat gtggcgtaat g 51

<210> 5915
<211> 51
<212> DNA

<213> Homo sapiens

<220>

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<222> (26)...(0)

<223> 1 of 2 allelic variants (5916 is other entry)

<221> misc_feature

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<223> Accession number cg44127585

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cgacctcgca ctcggcgatg gacaaaatgt ggcgtaatgg caagcgggta c

51

<210> 5916

<211> 51

<212> DNA

<213> Homo sapiens

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<222> (26)...(0)

<223> 2 of 2 allelic variants (5915 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44127585

<400> 5916

cgacctcgca ctcggcgatg gacaagatgt ggcgtaatgg caagcgggta c

51

<210> 5917

<211> 51

<212> DNA

<213> Homo sapiens

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<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (5918 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44127585

<400> 5917

atggcaagcg ggtacgacgg gtcaagctgg acgagaaccg caactgggaa a

51

<210> 5918

<211> 51

<212> DNA

<213> Homo sapiens

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<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (5917 is other entry)

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<223> Accession number cg44127585

<400> 5918
atggcaagcg ggtacgacgg gtcaaactgg acgagaaccg caactgggaa a 51

<210> 5919
<211> 51
<212> DNA
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<220>
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<223> 1 of 2 allelic variants (5920 is other entry)

<221> misc_feature
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<223> Accession number cg44127585

<400> 5919
gggtacgacg ggtcaagctg gacgagaacc gcaactggga aaagttccga g 51

<210> 5920
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<220>
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<223> 2 of 2 allelic variants (5919 is other entry)

<221> misc_feature
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<400> 5920
gggtacgacg ggtcaagctg gacgaaaacc gcaactggga aaagttccga g 51

<210> 5921
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<223> 1 of 2 allelic variants (5922 is other entry)

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ctggacgaga accgcaactg ggaaaagtgc cgagaggcca tgccggacgc t 51

<210> 5922

<211> 51

<212> DNA

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<223> 2 of 2 allelic variants (5921 is other entry)

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<400> 5922

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<210> 5923

<211> 51

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<223> 1 of 2 allelic variants (5924 is other entry)

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<222> (0)...(0)

<223> Accession number cg44127585

<400> 5923

agaaccgcaa ctgggaaaag ttccgagagg ccatgccgga cgctgtggtg a 51

<210> 5924

<211> 51

<212> DNA

<213> Homo sapiens

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<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (5923 is other entry)

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<223> Accession number cg44127585

<400> 5924

agaaccgcaa ctgggaaaag ttccgggagg ccatgccgga cgctgtggtg a 51

<210> 5925

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 1 of 2 allelic variants (5926 is other entry)

<221> misc_feature
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tgccggacgc tgtggtgatg tttgtcgta agccttcttc ggtgaacgtt g 51

<210> 5926
<211> 51
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<220>
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tgccggacgc tgtggtgatg tttgttgta agccttcttc ggtgaacgtt g 51

<210> 5927
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<223> 1 of 2 allelic variants (5928 is other entry)

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<400> 5927
gtgaacgttg ccaccaacgg gatgaatgtt cccaaagagg actacaccat c 51

<210> 5928
<211> 51
<212> DNA
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<223> 2 of 2 allelic variants (5927 is other entry)

<221> misc_feature
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<400> 5928
gtgaacgttg ccaccaacgg gatgagtgtt cccaaagagg actacaccat c 51

<210> 5929
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<223> 1 of 2 allelic variants (5930 is other entry)

<221> misc_feature
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<400> 5929
aacgttgcca ccaacgggat gaatgttccc aaagaggact acaccatcat t 51

<210> 5930
<211> 51
<212> DNA
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<223> 2 of 2 allelic variants (5929 is other entry)

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<222> (0)...(0)
<223> Accession number cg44127585

<400> 5930
aacgttgcca ccaacgggat gaatgttccc aaagaggact acaccatcat t 51

<210> 5931
<211> 51
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<223> 1 of 2 allelic variants (5932 is other entry)

<221> misc_feature
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<400> 5931
cgggatgaat gttcccaaag aggactacac catcatatat ctgggtgacg a 51

<210> 5932

<211> 51

<212> DNA

<213> Homo sapiens

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<222> (26)...(0)

<223> 2 of 2 allelic variants (5931 is other entry)

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<222> (0)...(0)

<223> Accession number cg44127585

<400> 5932

cgggatgaat gttcccaaag aggacgacac catcatttat ctgggtgacg a

51

<210> 5933

<211> 51

<212> DNA

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<223> 1 of 2 allelic variants (5934 is other entry)

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<223> Accession number cg44127585

<400> 5933

aagaggacta caccatcatt tatctgggtg acgagtggaa agacgctccg g

51

<210> 5934

<211> 51

<212> DNA

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<223> 2 of 2 allelic variants (5933 is other entry)

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51

<210> 5935

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 1 of 2 allelic variants (5936 is other entry)

<221> misc_feature
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<223> Accession number cg44128033

<400> 5935
accacttcag cgatctgcaa agcctggtgt cagcatccac ctccgagctc g 51

<210> 5936
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (5935 is other entry)

<221> misc_feature
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<223> Accession number cg44128033

<400> 5936
accacttcag cgatctgcaa agcctagtgt cagcatccac ctccgagctc g 51

<210> 5937
<211> 51
<212> DNA
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<220>
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<223> 1 of 2 allelic variants (5938 is other entry)

<221> misc_feature
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<223> Accession number cg44128033

<400> 5937
tgggtgtcagc atccacctcc gagctcggcg atattaaggg ggtcggcctc a 51

<210> 5938
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<223> 2 of 2 allelic variants (5937 is other entry)

<221> misc_feature

<222> (0)...(0)
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<400> 5938
tgggtgtcagc atccacctcc gagcttggcg atattaaggg ggtcggcctc a 51

<210> 5939
<211> 51
<212> DNA
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5940 is other entry)

<221> misc_feature
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<223> Accession number cg44128033

<400> 5939
gccggttgct cactgatgac tatcgtecca attccgtcgc tcctggcggg t 51

<210> 5940
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (5939 is other entry)

<221> misc_feature
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<400> 5940
gccggttgct cactgatgac tatcgcccca attccgtcgc tcctggcggg t 51

<210> 5941
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (5942 is other entry)

<221> misc_feature
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<223> Accession number cg44128033

<400> 5941
tcgctcctgg cgggttttcc ctgtcaggat tacagaaact gtcctgggaa g 51

<210> 5942
<211> 51
<212> DNA
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<220>
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<223> 2 of 2 allelic variants (5941 is other entry)

<221> misc_feature
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<223> Accession number cg44128033

<400> 5942
tcgctcctgg cgggttttcc ctgtccggat tacagaaact gtcctgggaa g 51

<210> 5943
<211> 51
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<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5944 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44128710

<400> 5943
gcgtgtgtga ctgtgaaggg gccgccggcg tctgtaggaa ggcacagcct g 51

<210> 5944
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5943 is other entry)

<221> misc_feature
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<223> Accession number cg44128710

<400> 5944
gcgtgtgtga ctgtgaaggg gccgctggcg tctgtaggaa ggcacagcct g 51

<210> 5945
<211> 51
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5946 is other entry)

<221> misc_feature
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<223> Accession number cg44128803

<400> 5945
taacttattg tatttttagt agagatggga tgtcaccatg ttggccagga t 51

<210> 5946
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (5945 is other entry)

<221> misc_feature
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<223> Accession number cg44128803

<400> 5946
taacttattg tatttttagt agagacggga tgtcaccatg ttggccagga t 51

<210> 5947
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5948 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44129401

<400> 5947
agggctgagt gccaaaagct tgggggagga gggacaggtc ctcggcaata a 51

<210> 5948
<211> 51
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<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (5947 is other entry)

<221> misc_feature
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<223> Accession number cg44129401

<400> 5948

agggctgagt gccaaaagct tggggaagga gggacaggtc ctcggcaata a

51

<210> 5949

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (5950 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44129401

<400> 5949

ctcggcaata aacagtgtca cacacgcac acagccattt gagaaatggc t

51

<210> 5950

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

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<223> 2 of 2 allelic variants (5949 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44129401

<400> 5950

ctcggcaata aacagtgtca cacacacac acagccattt gagaaatggc t

51

<210> 5951

<211> 50

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (5952 is other entry)

<221> misc_feature

<222> (25)...(26)

<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44129641

<400> 5951
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<210> 5952
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5951 is other entry)

<221> misc_feature
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<223> Accession number cg44129641

<400> 5952
gggtgcggtc tcctcgctgg cctcctcttc ttcacttggtg tcttcccatt a 51

<210> 5953
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (5954 is other entry)

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<223> Accession number cg44130064

<400> 5953
atttaacact attgtatttt tattatatgt aatttagtaa tatgaatata a 51

<210> 5954
<211> 51
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<213> Homo sapiens

<220>
<221> misc_feature
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<223> 2 of 2 allelic variants (5953 is other entry)

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<400> 5954
atttaacact attgtatttt tattacatgt aatttagtaa tatgaatata a 51

<210> 5955
<211> 51
<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (5956 is other entry)

<221> misc_feature

<222> (0)...(0)

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gtacactggt gtttatatatt gcacagagta ttgatatgtg atgtattaag t

51

<210> 5956

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<212> DNA

<213> Homo sapiens

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<223> 2 of 2 allelic variants (5955 is other entry)

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51

<210> 5957

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<212> DNA

<213> Homo sapiens

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<223> 1 of 2 allelic variants (5958 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44130288

<400> 5957

tagccagatg taaccagctt gctgtcttgt cccaagcctc cctctaaggg g

51

<210> 5958

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (5957 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44130288

<400> 5958
tagccagatg taaccagctt gctgttttgt cccaagcctc cctctaagg g 51

<210> 5959
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5960 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44130288

<400> 5959
cttgtcccaa gcctccctct aaggggacag tgtgaatcgg tgaatgttga g 51

<210> 5960
<211> 50
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (5959 is other entry)

<221> misc_feature
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<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44130288

<400> 5960
cttgtcccaa gcctccctct aagggacagt gtgaatcggg gaatgttgag 50

<210> 5961
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5962 is other entry)

<221> misc_feature

<222> (0)...(0)
<223> Accession number cg44130288

<400> 5961
gggaatgctc tgaaatcagt gtgggcatgg ctgtaccaac aggaatgaac a 51

<210> 5962
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (5961 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44130288

<400> 5962
gggaatgctc tgaaatcagt gtgggaatgg ctgtaccaac aggaatgaac a 51

<210> 5963
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (5964 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44130288

<400> 5963
ccaacaggaa tgaacagttg ttcgatgaca gatgactccc aagtgcaca c 51

<210> 5964
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (5963 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44130288

<400> 5964
ccaacaggaa tgaacagttg ttcgacgaca gatgactccc aagtgcaca c 51

<210> 5965
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5966 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44130288

<400> 5965
atgacagatg actcccaagt gacacacacg tgcctcaag aagactcaca c 51

<210> 5966
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5965 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44130288

<400> 5966
atgacagatg actcccaagt gacacgcacg tgcctcaag aagactcaca c 51

<210> 5967
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5968 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44130564

<400> 5967
aacctgatca agcaggatga cggcggtcc cccatcagac actatctggt c 51

<210> 5968
<211> 51
<212> DNA
<213> Homo sapiens

<220>

<221> misc_feature
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<223> 2 of 2 allelic variants (5967 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44130564

<400> 5968
aacctgatca agcaggatga cggcgactcc cccatcagac actatctggt c 51

<210> 5969
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5970 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44130564

<400> 5969
cctcaccatc tataacgcca acatcgacga cgccggcatt tacaagtgtg t 51

<210> 5970
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (5969 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44130564

<400> 5970
cctcaccatc tataacgcca acatctacga cgccggcatt tacaagtgtg t 51

<210> 5971
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5972 is other entry)

<221> misc_feature
<222> (0)...(0)

<223> Accession number cg44130566

<400> 5971

cgtgtgtgtg tgtgtgtgtg tgtgtgtgtt ttaattttta ttttagagac a

51

<210> 5972

<211> 50

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (5971 is other entry)

<221> misc_feature

<222> (25)...(26)

<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44130566

<400> 5972

cgtgtgtgtg tgtgtgtgtg tgtgtgttt taatttttat ttttagagaca

50

<210> 5973

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (5974 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44130566

<400> 5973

tgtgtgtgtg tgtgtgtgtg tgtgtgttt aatttttatt ttagagacag g

51

<210> 5974

<211> 50

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (5973 is other entry)

<221> misc_feature

<222> (25)...(26)

<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44130566

<400> 5974
tgtgtgtgtg tgtgtgtgtg tgtgttttta atttttattt tagagacagg 50

<210> 5975
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5976 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44130641

<400> 5975
agttcccaaa cttagggtgc ccaatgtccc tcaccttgaa gttagcaaga g 51

<210> 5976
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (5975 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44130641

<400> 5976
agttcccaaa cttagggtgc ccaatatccc tcaccttgaa gttagcaaga g 51

<210> 5977
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5978 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44131199

<400> 5977
agcaggacgt cgctgccggg ggagccctcc agacacgcag aaatgccaca g 51

<210> 5978
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5977 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44131199

<400> 5978
agcaggacgt cgctgccggg ggagctotcc agacacgcag aaatgccaca g 51

<210> 5979
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5980 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44131207

<400> 5979
gaattcggtg cgtttggtgg ctatggcacc ctcaccagct ttgacatcca t 51

<210> 5980
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (5979 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44131207

<400> 5980
gaattcggtg cgtttggtgg ctatgacacc ctcaccagct ttgacatcca t 51

<210> 5981
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5982 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44131588

<400> 5981
agctttctcac ctctacggtg aaagtttcgg tgtcacccac atggggcttt t 51

<210> 5982
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (5981 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44131588

<400> 5982
agctttctcac ctctacggtg aaagtctcgg tgtcacccac atggggcttt t 51

<210> 5983
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5984 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44131588

<400> 5983
tgcgaaccct cacacatcgc tgacgacggc ggcggagtcg atgctgaggc c 51

<210> 5984
<211> 50
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5983 is other entry)

<221> misc_feature

<222> (25)...(26)
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<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44131588

<400> 5984
tgccaaccct,cacacatcgc tgacgcggcg gcggagtcga tgctgaggcc 50

<210> 5985
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5986 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44131588

<400> 5985
ctgacgcgcg cggcggagtc gatgctgagg ccgacggcga tggggatttc g 51

<210> 5986
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (5985 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44131588

<400> 5986
ctgacgcgcg cggcggagtc gatgccgagg ccgacggcga tggggatttc g 51

<210> 5987
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5988 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44131588

<400> 5987
gctgaggccg acggcgatgg ggatttcgtc cggggcgag acctacttgg g 51

<210> 5988
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5987 is other entry)

<221> misc_feature
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<223> Accession number cg44131588

<400> 5988
gctgaggccg acggcgatgg ggatttcgtc cggggcgag acctacttgg g 51

<210> 5989
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5990 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44131701

<400> 5989
gcggtgccag aaaaaggcca aataactctgc tttcccgagg gacgcgcgta g 51

<210> 5990
<211> 50
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (5989 is other entry)

<221> misc_feature
<222> (25)...(26)
<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44131701

<400> 5990

gcggtgccag aaaaaggcca aatacctgct ttcccgaggg acgcgcgtag

50

<210> 5991

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (5992 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44131756

<400> 5991

acagcagggc catgggaggg gggcccggt cactcggcgg agcccttggtg g

51

<210> 5992

<211> 50

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (5991 is other entry)

<221> misc_feature

<222> (25)...(26)

<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44131756

<400> 5992

acagcagggc catgggaggg gggccggctc actcggcgga gcccttggtg

50

<210> 5993

<211> 51

<212> DNA

<213> Homo sapiens

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<222> (26)...(0)

<223> 1 of 2 allelic variants (5994 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44131756

<400> 5993

aggcattgag cggagatctc accactgcac tccagcctgg gcgacagagc a

51

<210> 5994
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (5993 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44131756

<400> 5994
aggcattgag cggagatctc accaccgcac tccagcctgg ggcacagagc a 51

<210> 5995
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (5996 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44131756

<400> 5995
acagagcaag attctatcaa aaaagaaaga aagaaaggac aggacaggaa a 51

<210> 5996
<211> 50
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (5995 is other entry)

<221> misc_feature
<222> (25)...(26)
<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44131756

<400> 5996
acagagcaag attctatcaa aaaagaagaa agaaaggaca ggacaggaaa 50

<210> 5997
<211> 51

<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (5998 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44131756

<400> 5997
agattctatc aaaaaagaaa gaaagaaagg acaggacagg aaaggaagg g 51

<210> 5998
<211> 50
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (5997 is other entry)

<221> misc_feature
<222> (25)...(26)
<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44131756

<400> 5998
agattctatc aaaaaagaaa gaaagaagga caggacagga aaggaaggg 50

<210> 5999
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (6000 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44131756

<400> 5999
gattctatca aaaaagaaag aaagaaagga caggacagga aaggaaggg g 51

<210> 6000
<211> 50
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (5999 is other entry)

<221> misc_feature
<222> (25)...(26)
<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44131756

<400> 6000
gattctatca aaaagaaag aaagaaggac aggacaggaa agggaagggg 50

<210> 6001
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (6002 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44131756

<400> 6001
attctatcaa aaaagaaaga aagaaggac aggacaggaa agggaagggg t 51

<210> 6002
<211> 50
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (6001 is other entry)

<221> misc_feature
<222> (25)...(26)
<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44131756

<400> 6002
attctatcaa aaaagaaaga aagaaggaca ggacaggaaa ggggaaggggt 50

<210> 6003
<211> 50

<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (6004 is other entry)

<221> misc_feature
<222> (25)...(26)
<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44131756

<400> 6003
cagaactggg ggcaaaaaca aaaaagaagg aaggaaggaa agaaagaaat 50

<210> 6004
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (6003 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44131756

<400> 6004
cagaactggg ggcaaaaaca aaaaagaag gaaggaagga aagaagaaa t 51

<210> 6005
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (6006 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44131756

<400> 6005
aagaaggaag gaaggaaaga aagaaatggg tatgggcaga attgggggca a 51

<210> 6006
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (6005 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44131756

<400> 6006
aagaaggaag gaaggaaaga aagaagtggg tatgggcaga attgggggca a 51

<210> 6007
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (6008 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44910941

<400> 6007
gggccagact gagccatgcc acacccttcc tcctagtccc catgctctcc t 51

<210> 6008
<211> 50
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (6007 is other entry)

<221> misc_feature
<222> (25)...(26)
<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44910941

<400> 6008
gggccagact gagccatgcc acacccttcc cctagtcccc atgctctcct 50

<210> 6009
<211> 51
<212> DNA
<213> Homo sapiens

<220>

<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (6010 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44911060

<400> 6009
tggaatggaa tgcaatggaa tggaatcgac tggaatggaa tggaatggaa t 51

<210> 6010
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (6009 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44911060

<400> 6010
tggaatggaa tgcaatggaa tggaaccgac tggaatggaa tggaatggaa t 51

<210> 6011
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (6012 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44911060

<400> 6011
ggaatggaat gcaatggaat ggaatcgact ggaatggaat ggaatggaat g 51

<210> 6012
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
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<223> 2 of 2 allelic variants (6011 is other entry)

<221> misc_feature
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<223> Accession number cg44911060

<400> 6012

ggaatggaat gcaatggaat ggaatggact ggaatggaat ggaatggaat g 51

<210> 6013

<211> 51

<212> DNA

<213> Homo sapiens

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<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (6014 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44911060

<400> 6013

ggaatggaat ggaatggaat ggaatcaacc cgagtgcaat ggaatggagt g 51

<210> 6014

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 2 of 2 allelic variants (6013 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44911060

<400> 6014

ggaatggaat ggaatggaat ggaataaacc cgagtgcaat ggaatggagt g 51

<210> 6015

<211> 51

<212> DNA

<213> Homo sapiens

<220>

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<222> (26)...(0)

<223> 1 of 2 allelic variants (6016 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44911343

<400> 6015

tttttttttt ttgaaacagg gtctcactct gctgccacagg ctggagtgca g 51

<210> 6016

<211> 51
<212> DNA
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<220>
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<223> 2 of 2 allelic variants (6015 is other entry)

<221> misc_feature
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<223> Accession number cg44911343

<400> 6016
tttttttttt ttgaaacagg gtctcgctct gctgcccagg ctggagtga g 51

<210> 6017
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (6018 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44911865

<400> 6017
accgttgct agataccagg ctgcgccatt ccaatcccg atccgcgtca c 51

<210> 6018
<211> 50
<212> DNA
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<220>
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<221> misc_feature
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<221> misc_feature
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<223> Accession number cg44911865

<400> 6018
accgttgct agataccagg ctgcgcattc caatcccgga tccgcgtcac 50

<210> 6019
<211> 51
<212> DNA

<213> Homo sapiens

<220>

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<223> 1 of 2 allelic variants (6020 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44911865

<400> 6019

ccgttgcgta gataccaggc tgcgccattc caatcccgga tccgcgtcac a

51

<210> 6020

<211> 50

<212> DNA

<213> Homo sapiens

<220>

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<222> (26)...(0)

<223> 2 of 2 allelic variants (6019 is other entry)

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<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44911865

<400> 6020

ccgttgcgta gataccaggc tgcgcattcc aatcccggat ccgcgtcaca

50

<210> 6021

<211> 46

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

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<223> 1 of 2 allelic variants (6022 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44911865

<400> 6021

gtcgatcgat gtcgcgcgcg gtcgcctctg cgaggatagc acgcgt

46

<210> 6022

<211> 45

<212> DNA

<213> Homo sapiens

<220> .
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<223> 2 of 2 allelic variants (6021 is other entry)

<221> misc_feature
<222> (25)...(26)
<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44911865

<400> 6022
gtcgatcgat gctcgcgccg gtcgctctgc gaggatacga cgcgt

45

<210> 6023
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (6024 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44912335

<400> 6023
accctgagtc atgggggtct tttgtaaaag tccccaaact ggcccaggaa g

51

<210> 6024
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (6023 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44912335

<400> 6024
accctgagtc atgggggtct tttgtgaaag tccccaaact ggcccaggaa g

51

<210> 6025
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (6026 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44912347

<400> 6025
ctctaattga agctctggca tcattctgggg ctttatgagc caagggagat a 51

<210> 6026
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (6025 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44912347

<400> 6026
ctctaattga agctctggca tcattcagggg ctttatgagc caagggagat a 51

<210> 6027
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (6028 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44913034

<400> 6027
gtcagggtgc tcggcaaggg cgagcatcct ccatgcagca gagacaccag a 51

<210> 6028
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (6027 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44913034

<400> 6028
gtcaggggtgc tcggcaaggg cgagcgtcct ccatgcagca gagacaccag a 51

<210> 6029
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (6030 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44913283

<400> 6029
gtttggcact cgctcatgaat tgaagaatga aaagccatag tcacaagtct g 51

<210> 6030
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (6029 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44913283

<400> 6030
gtttggcact cgctcatgaat tgaaggatga aaagccatag tcacaagtct g 51

<210> 6031
<211> 51
<212> DNA
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<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (6032 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44913564

<400> 6031
cggccgtgaa gagcagcctg gcggacggtg taaacctgca caggtgcact g 51

<210> 6032
<211> 51

<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (6031 is other entry)

<221> misc_feature
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<223> Accession number cg44913564

<400> 6032
cggccgtgaa gaggcgcctg gcggaagggtg taaacctgca caggtgcact g 51

<210> 6033
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (6034 is other entry)

<221> misc_feature
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<223> Accession number cg44913901

<400> 6033
aggcggagggt tgcagtgagc cgagaccaca ccattgcact ccagcctgga c 51

<210> 6034
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (6033 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44913901

<400> 6034
aggcggagggt tgcagtgagc cgagatcaca ccattgcact ccagcctgga c 51

<210> 6035
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (6036 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44914000

<400> 6035

aactatntag gttgtttcta attttgatta ttataaagtt gcagaaattt g

51

<210> 6036

<211> 51

<212> DNA

<213> Homo sapiens

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<222> (26)...(0)

<223> 2 of 2 allelic variants (6035 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44914000

<400> 6036

aactatntag gttgtttcta attttcatta ttataaagtt gcagaaattt g

51

<210> 6037

<211> 51

<212> DNA

<213> Homo sapiens

<220>

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<222> (26)...(0)

<223> 1 of 2 allelic variants (6038 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44914031

<400> 6037

ggtgacagag tgagacactg tctccaaaaa aaaaatgttt aaaatgagac c

51

<210> 6038

<211> 50

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (6037 is other entry)

<221> misc_feature

<222> (25)...(26)

<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44914031

<400> 6038
ggtgacagag tgagacactg tctccaaaaa aaaatgttta aaatgagacc

50

<210> 6039
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (6040 is other entry)

<221> misc_feature
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<223> Accession number cg44914031

<400> 6039
caaaagattc tacttccttg cttgggcggg gctccgattc tccaaactga t

51

<210> 6040
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (6039 is other entry)

<221> misc_feature
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<223> Accession number cg44914031

<400> 6040
caaaagattc tacttccttg cttggacggg gctccgattc tccaaactga t

51

<210> 6041
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (6042 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44914076

<400> 6041
tgagggtcaag ctgtctgccc atctcagcct cccaaagtgc tgggattgca g

51

<210> 6042
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (6041 is other entry)

<221> misc_feature
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<223> Accession number cg44914076

<400> 6042
tgaggtcaag ctgtctgccc atctcgccct cccaaagtgc tgggattgca g 51

<210> 6043
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (6044 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44914681

<400> 6043
cttctagaca caccctgagc cagaaggga cccactgcct tgaagggaag g 51

<210> 6044
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (6043 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44914681

<400> 6044
cttctagaca caccctgagc cagaatggaa cccactgcct tgaagggaag g 51

<210> 6045
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (6046 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44914784

<400> 6045
tccagctgga gcggatggat gagagtgatg acggtgacgc cactaccggc a 51

<210> 6046
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (6045 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44914784

<400> 6046
tccagctgga gcggatggat gagagagatg acggtgacgc cactaccggc a 51

<210> 6047
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (6048 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44914784

<400> 6047
cgacctcgca ctcggcgatg gacaaaatgt ggcgtaatgg caagcgggta c 51

<210> 6048
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (6047 is other entry)

<221> misc_feature

<222> (0)...(0)
<223> Accession number cg44914784

<400> 6048
cgacctcgca ctcggcgatg gacaagatgt ggcgtaatgg caagcgggta c 51

<210> 6049
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (6050 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44914784

<400> 6049
atggcaagcg ggtacgacgg gtcaagctgg acgagaaccg caactgggaa a 51

<210> 6050
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (6049 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44914784

<400> 6050
atggcaagcg ggtacgacgg gtcaaaactgg acgagaaccg caactgggaa a 51

<210> 6051
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (6052 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44914784

<400> 6051
gggtacgacg ggtcaagctg gacgagaacc gcaactggga aaagttccga g 51

<210> 6052
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (6051 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44914784

<400> 6052
gggtacgacg ggtcaagctg gacgaaaacc gcaactggga aaagttccga g 51

<210> 6053
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (6054 is other entry)

<221> misc_feature
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<223> Accession number cg44914784

<400> 6053
ctggacgaga accgcaactg ggaaaagtgc cgagaggcca tgccggacgc t 51

<210> 6054
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (6053 is other entry)

<221> misc_feature
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<223> Accession number cg44914784

<400> 6054
ctggacgaga accgcaactg ggaaatgttc cgagaggcca tgccggacgc t 51

<210> 6055
<211> 51
<212> DNA
<213> Homo sapiens

<220>

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<223> 1 of 2 allelic variants (6056 is other entry)

<221> misc_feature
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<223> Accession number cg44914784

<400> 6055
agaaccgcaa ctgggaaaag ttccgagagg ccatgccgga cgctgtggtg a 51

<210> 6056
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (6055 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44914784

<400> 6056
agaaccgcaa ctgggaaaag ttccgggagg ccatgccgga cgctgtggtg a 51

<210> 6057
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (6058 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44914811

<400> 6057
cgtgagcaat gacgtcagcc atagtgggtga ggcggtcatc gctgaccccg g 51

<210> 6058
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (6057 is other entry)

<221> misc_feature
<222> (0)...(0)

<223> Accession number cg44914811

<400> 6058
cgtgagcaat gacgtcagcc atagtagtga ggcggtcac gctgaccccg g 51

<210> 6059
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (6060 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44914811

<400> 6059
tgaccccggt ctgctcggag tcgtcgacaa agctaataatgt gtagcgtacc a 51

<210> 6060
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (6059 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44914811

<400> 6060
tgaccccggt ctgctcggag tcgtcaacaa agctaataatgt gtagcgtacc a 51

<210> 6061
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (6062 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44914811

<400> 6061
tgctcggagt cgtcgacaaa gctaataatgt gtagcgtacca ccgcgtcgac t 51

<210> 6062

<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (6061 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44914811

<400> 6062
tgctcggagt cgtcgacaaa gctaacgatg tgacgtacca ccgcgtcgac t 51

<210> 6063
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (6064 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44914811

<400> 6063
aattcgggta gatcgtcgta gtcgcaccac ctgcgcagga aatggccata t 51

<210> 6064
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (6063 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44914811

<400> 6064
aattcgggta gatcgtcgta gtcgcgccac ctgcgcagga aatggccata t 51

<210> 6065
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature

<222> (26)...(0)
<223> 1 of 2 allelic variants (6066 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44914838

<400> 6065
tggcagtcgg cggacagacg ccgctagcat cagggtcttc atgcgctgga a 51

<210> 6066
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (6065 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44914838

<400> 6066
tggcagtcgg cggacagacg ccgctggcat cagggtcttc atgcgctgga a 51

<210> 6067
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (6068 is other entry)

<221> misc_feature
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<223> Accession number cg44914838

<400> 6067
ttgacggtcg ggtggaccga actgctacca atcaggcatt cgatgctgca c 51

<210> 6068
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (6067 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44914838

<400> 6068
ttgacggtcg ggtggaccga actgccacca atcaggcatt cgatgtcgca c 51

<210> 6069
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (6070 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44914838

<400> 6069
atcaggcatt cgatgtcgca caaaccttgt ccgacgacac aagaacctcg t 51

<210> 6070
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (6069 is other entry)

<221> misc_feature
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<223> Accession number cg44914838

<400> 6070
atcaggcatt cgatgtcgca caaaccttgt ccgacgacac aagaacctcg t 51

<210> 6071
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (6072 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44914838

<400> 6071
tcgtagatac ctccgtcaac cgagttcatc aaagggactg cgaagcgctg t 51

<210> 6072
<211> 51

<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (6071 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44914838

<400> 6072
tcgtagatac ctccgtcaac cgagtccatc aaagggactg cgaagcgtg t 51

<210> 6073
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (6074 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44914838

<400> 6073
gagttcatca aagggactgc gaagcgtgt caggaattct tcatgcgcct c 51

<210> 6074
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (6073 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44914838

<400> 6074
gagttcatca aagggactgc gaagcactgt caggaattct tcatgcgcct c 51

<210> 6075
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (6076 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44914862

<400> 6075

gccccgtact cacctcagac atagttcacc catctcttct gccccaggaa c

51

<210> 6076

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (6075 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44914862

<400> 6076

gccccgtact cacctcagac atagttcacc catctcttct gccccaggaa c

51

<210> 6077

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (6078 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44914862

<400> 6077

actcacctca gacatagttc acccatctct tctgccccag gaactggacg a

51

<210> 6078

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (6077 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44914862

<400> 6078
actcacctca gacatagttc acccacctct tctgccccag gaactggacg a 51

<210> 6079
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (6080 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44914862

<400> 6079
ccatctcttc tgccccagga actggacgag ggtcacgatg gcaacgaggg c 51

<210> 6080
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (6079 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44914862

<400> 6080
ccatctcttc tgccccagga actgggcgag ggtcacgatg gcaacgaggg c 51

<210> 6081
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (6082 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44914862

<400> 6081
tcttctgccc caggaactgg acgaggggtca cgatggcaac gagggcaaag a 51

<210> 6082
<211> 51
<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (6081 is other entry)

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<222> (0)...(0)

<223> Accession number cg44914862

<400> 6082

tcttctgccc caggaactgg acgagagtca cgatggcaac gagggcaaag a

51

<210> 6083

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (6084 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44914862

<400> 6083

ggaactggac gagggtcacg atggcaacga gggcaaagat aacgatcgcc a

51

<210> 6084

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (6083 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44914862

<400> 6084

ggaactggac gagggtcacg atggcgacga gggcaaagat aacgatcgcc a

51

<210> 6085

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (6086 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44914862

<400> 6085
ggacgagggt cacgatggca acgagggcaa agataacgat cgccacggca g 51

<210> 6086
<211> 51
<212> DNA
<213> Homo sapiens

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<222> (26)...(0)
<223> 2 of 2 allelic variants (6085 is other entry)

<221> misc_feature
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<223> Accession number cg44914862

<400> 6086
ggacgagggt cacgatggca acgagagcaa agataacgat cgccacggca g 51

<210> 6087
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (6088 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44914862

<400> 6087
gggcaaagat aacgatcgcc acggcagcac cgagcccctg gtcatggttg a 51

<210> 6088
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 2 of 2 allelic variants (6087 is other entry)

<221> misc_feature
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<223> Accession number cg44914862

<400> 6088

gggcaaagat aacgatcgcc acggcggcac cgagcccctg gtcatggttg a

51

<210> 6089

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (6090 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44914862

<400> 6089

cctggtcatg gttgacgaag gcttggcggg agaacagcga caccaacgag c

51

<210> 6090

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (6089 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44914862

<400> 6090

cctggtcatg gttgacgaag gcttgtcggg agaacagcga caccaacgag c

51

<210> 6091

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (6092 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44914862

<400> 6091

gcttggcggg agaacagcga caccaacgag cgagtcctcg gctcagccgg g

51

<210> 6092

<211> 51

<212> DNA

<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (6091 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44914862

<400> 6092
gcttggcggg agaacagcga caccagcgag cgagtcctcg gctcagccgg g 51

<210> 6093
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (6094 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44914862

<400> 6093
ttccgatcat cgcgtacagc gcgtcaaaca actggaaccc accgatcgtc g 51

<210> 6094
<211> 51
<212> DNA
<213> Homo sapiens

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<222> (26)...(0)
<223> 2 of 2 allelic variants (6093 is other entry)

<221> misc_feature
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<223> Accession number cg44914862

<400> 6094
ttccgatcat cgcgtacagc gcgtcgaaca actggaaccc accgatcgtc g 51

<210> 6095
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (6096 is other entry)

<221> misc_feature
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<223> Accession number cg44914862

<400> 6095
acgaagcgcc gctgcgacac ggtgtcgccc cgcttggcaa gggccgccac c 51

<210> 6096
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (6095 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44914862

<400> 6096
acgaagcgcc gctgcgacac ggtgttgccc cgcttggcaa gggccgccac c 51

<210> 6097
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (6098 is other entry)

<221> misc_feature
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<223> Accession number cg44914862

<400> 6097
cggtgtcgcc ccgcttggca agggccgcca cctctacagc ggagagttcg g 51

<210> 6098
<211> 51
<212> DNA
<213> Homo sapiens

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<222> (26)...(0)
<223> 2 of 2 allelic variants (6097 is other entry)

<221> misc_feature
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<223> Accession number cg44914862

<400> 6098
cggtgtcgcc ccgcttggca agggctgcca cctctacagc ggagagttcg g 51

<210> 6099
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (6100 is other entry)

<221> misc_feature
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<223> Accession number cg44914862

<400> 6099
gccacctcta cagcggagag ttcgggcccc gtactcacct cagacatagt t 51

<210> 6100
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (6099 is other entry)

<221> misc_feature
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<400> 6100
gccacctcta cagcggagag ttcgggcccc gtactcacct cagacatagt t 51

<210> 6101
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (6102 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44914862

<400> 6101
tacagcggag agttcgggcc ccgtactcac ctcagacata gttcacccat c 51

<210> 6102
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (6101 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44914862

<400> 6102
tacagcggag agttcgggcc ccgtaatcac ctcagacata gttcacccat c 51

<210> 6103
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (6104 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44914864

<400> 6103
cctcaagatc agccgtacgg ccggtaggga ccgcactaac cttcgggcct c 51

<210> 6104
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (6103 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44914864

<400> 6104
cctcaagatc agccgtacgg ccggtgagga ccgcactaac cttcgggcct c 51

<210> 6105
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (6106 is other entry)

<221> misc_feature

<222> (0)...(0)
<223> Accession number cg44914864

<400> 6105
ccttcgggcc tcggtcgaag tcggcgcgcg agcctgatcc gaagaaggac g 51

<210> 6106
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (6105 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44914864

<400> 6106
ccttcgggcc tcggtcgaag tcggcacgcg agcctgatcc gaagaaggac g 51

<210> 6107
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (6108 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44914864

<400> 6107
ggcctcggtc gaagtcggcg cgcgagcctg atccgaagaa ggacgaggtg g 51

<210> 6108
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (6107 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44914864

<400> 6108
ggcctcggtc gaagtcggcg cgcgatcctg atccgaagaa ggacgaggtg g 51

<210> 6109
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (6110 is other entry)

<221> misc_feature
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<223> Accession number cg44914864

<400> 6109
ggtcgaagtc ggcgcgcgag cctgatccga agaaggacga ggtgggagcc c 51

<210> 6110
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (6109 is other entry)

<221> misc_feature
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<223> Accession number cg44914864

<400> 6110
ggtcgaagtc ggcgcgcgag cctgaccga agaaggacga ggtgggagcc c 51

<210> 6111
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (6112 is other entry)

<221> misc_feature
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<223> Accession number cg44914864

<400> 6111
cggcgcgcga gcctgatccg aagaaggacg aggtgggagc cccgaaaccc t 51

<210> 6112
<211> 51
<212> DNA
<213> Homo sapiens

<220>

<221> misc_feature
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<223> 2 of 2 allelic variants (6111 is other entry)

<221> misc_feature
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<223> Accession number cg44914864

<400> 6112
cggcgcgcgga gcctgatccg aagaaagacg aggtgggagc cccgaaaccc t 51

<210> 6113
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (6114 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44914864

<400> 6113
acgaggtggg agccccgaaa ccctcgcttt gagccgccac gcgagcagcg a 51

<210> 6114
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (6113 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44914864

<400> 6114
acgaggtggg agccccgaaa ccctcacttt gagccgccac gcgagcagcg a 51

<210> 6115
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (6116 is other entry)

<221> misc_feature
<222> (0)...(0)

<223> Accession number cg44914864

<400> 6115

tctcgacgac cgctgttggtg accgttaggg ttccacgact agccttctca c

51

<210> 6116

<211> 51

<212> DNA

<213> Homo sapiens

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<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (6115 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44914864

<400> 6116

tctcgacgac cgctgttggtg accgtcaggg ttccacgact agccttctca c

51

<210> 6117

<211> 51

<212> DNA

<213> Homo sapiens

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<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (6118 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44914864

<400> 6117

acgaccgctg ttgtgaccgt taggggtcca cgactagcct tctcacgatt g

51

<210> 6118

<211> 51

<212> DNA

<213> Homo sapiens

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<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (6117 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44914864

<400> 6118

acgaccgctg ttgtgaccgt taggggtcca cgactagcct tctcacgatt g

51

<210> 6119

<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (6120 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44914866

<400> 6119
gcccgctggt aatagcacga gctagctgga tggctgccga cccgacgcca c 51

<210> 6120
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (6119 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44914866

<400> 6120
gcccgctggt aatagcacga gctagtggga tggctgccga cccgacgcca c 51

<210> 6121
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (6122 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44914870

<400> 6121
tcctgtggag acgtggagtt ttccgcgcgt acctgggaga ttatttcgcc a 51

<210> 6122
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature

<222> (26)...(0)
<223> 2 of 2 allelic variants (6121 is other entry)

<221> misc_feature
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<223> Accession number cg44914870

<400> 6122
tcctgtggag acgtggagtt ttccgtgcgt acctgggaga ttatttcgcc a 51

<210> 6123
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (6124 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44914873

<400> 6123
ggcatgaacg accagcttct tgttactggc tagcgtcctg atcctcccag a 51

<210> 6124
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (6123 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44914873

<400> 6124
ggcatgaacg accagcttct tgttattggc tagcgtcctg atcctcccag a 51

<210> 6125
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (6126 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44914940

<400> 6125
cgcgatgtgc tgctcgctga ccgggaaact ctgacgaaga cggctagggt g 51

<210> 6126
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (6125 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44914940

<400> 6126
cgcgatgtgc tgctcgctga ccggggaact ctgacgaaga cggctagggt g 51

<210> 6127
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (6128 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44914951

<400> 6127
aaattggtga cctcaaggcc gatgtcaaac atggctgagc cggctcgag a 51

<210> 6128
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (6127 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44914951

<400> 6128
aaattggtga cctcaaggcc gatgttaaac atggctgagc cggctcgag a 51

<210> 6129
<211> 51

<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (6130 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44914951

<400> 6129
cgcagagtta cagtgggtcg cgcaaaataa tgggtgccga aacctacgtg g 51

<210> 6130
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (6129 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44914951

<400> 6130
cgcagagtta cagtgggtcg cgcaacataa tgggtgccga aacctacgtg g 51

<210> 6131
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (6132 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44914951

<400> 6131
aaaataatgg tgcccgaac ctacgtggta ctcatattca ctggacaatg t 51

<210> 6132
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (6131 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44914951

<400> 6132

aaaataatgg tgcccgaac ctacgcggta ctcattattca ctggacaatg t

51

<210> 6133

<211> 46

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (6134 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44914951

<400> 6133

cagggttcgt gccttgtaac aagacttgcc gagcgcatg gtgcac

46

<210> 6134

<211> 46

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (6133 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44914951

<400> 6134

cagggttcgt gccttgtaac aagacgtgcc gagcgcatg gtgcac

46

<210> 6135

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (6136 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44914955

<400> 6135
caaaccgcta cgagaccgcc tgcgaccac gtctgaaccg gaaccagtcc t 51

<210> 6136
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (6135 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44914955

<400> 6136
caaaccgcta cgagaccgcc tgcgatccac gtctgaaccg gaaccagtcc t 51

<210> 6137
<211> 43
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (18)...(0)
<223> 1 of 2 allelic variants (6138 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44914955

<400> 6137
gtgcacatcg agttgaccgg ggacgacgtc acggagtgtc tgg 43

<210> 6138
<211> 43
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (18)...(0)
<223> 2 of 2 allelic variants (6137 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44914955

<400> 6138
gtgcacatcg agttgactgg ggacgacgtc acggagtgtc tgg 43

<210> 6139
<211> 49
<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (6140 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44914955

<400> 6139

gtcagtcgcg tggcaagggtt ccgcagcatc gacctatgat cgatccgga

49

<210> 6140

<211> 49

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (6139 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44914955

<400> 6140

gtcagtcgcg tggcaagggtt ccgcaacatc gacctatgat cgatccgga

49

<210> 6141

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (6142 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44914955

<400> 6141

acgacgtcac ggagtgcttg ggcggggctcg acaagctagc agagtccgat c

51

<210> 6142

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (6141 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44914955

<400> 6142
acgacgtcac ggagtgcttg ggcggcgtcg acaagctagc agagtccgat c 51

<210> 6143
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (6144 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44914955

<400> 6143
agtgtcttggg cggggtcgac aagctagcag agtccgatct gacaaaccgc t 51

<210> 6144
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (6143 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44914955

<400> 6144
agtgtcttggg cggggtcgac aagctcgag agtccgatct gacaaaccgc t 51

<210> 6145
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (6146 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44914955

<400> 6145

gatctgacaa accgctacga gaccgcctgc gacccacgtc tgaaccggaa c 51

<210> 6146

<211> 51

<212> DNA

<213> Homo sapiens

<220>

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<222> (26)...(0)

<223> 2 of 2 allelic variants (6145 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44914955

<400> 6146

gatctgacaa accgctacga gaccgtctgc gacccacgtc tgaaccggaa c 51

<210> 6147

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (6148 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44914955

<400> 6147

tgacaaaccg ctacgagacc gcctgcgacc cacgtctgaa ccggaaccag t 51

<210> 6148

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (6147 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44914955

<400> 6148

tgacaaaccg ctacgagacc gcctgtgacc cacgtctgaa ccggaaccag t 51

<210> 6149

<211> 51

<212> DNA

<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (6150 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44914965

<400> 6149
ggctacaagt ttcaggccta aagcatcacc acccaggaaa gcaacaccac g 51

<210> 6150
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (6149 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44914965

<400> 6150
ggctacaagt ttcaggccta aagcaacacc acccaggaaa gcaacaccac g 51

<210> 6151
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (6152 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44914965

<400> 6151
taaagcatca ccacccagga aagcaacacc acgcaggaat ggtgtattca g 51

<210> 6152
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (6151 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44914965

<400> 6152
taaagcatca ccaccagga aagcagcacc acgcaggaat ggtgtattca g 51

<210> 6153
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (6154 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44914965

<400> 6153
tcagctgatt aggaccgagc cgacgaaatt cctgaagaag cgtactccgc a 51

<210> 6154
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (6153 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44914965

<400> 6154
tcagctgatt aggaccgagc cgacggaatt cctgaagaag cgtactccgc a 51

<210> 6155
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (6156 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44915044

<400> 6155
accgcgtcca agagctcgcg gatatggtct ctgaaaagta tgccgatgct g 51

<210> 6156
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (6155 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44915044

<400> 6156
accgctcca agagctcgcg gatatagtct ctgaaaagta tgccgatgct g 51

<210> 6157
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (6158 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44915044

<400> 6157
atatggtctc tgaaaagtat gccgatgctg agcaagacct gcttttggtc t 51

<210> 6158
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (6157 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44915044

<400> 6158
atatggtctc tgaaaagtat gccgacgctg agcaagacct gcttttggtc t 51

<210> 6159
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (6160 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44915044

<400> 6159
tgcttttggg ctgcgtgctc aagggcgcgg ctttcttctt gacggacttc g 51

<210> 6160
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (6159 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44915044

<400> 6160
tgcttttggg ctgcgtgctc aagggcgcgg ctttcttctt gacggacttc g 51

<210> 6161
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (6162 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44915044

<400> 6161
tgctcaaggg cgcggtcttc ttcttgacgg acttcgcgcg caagctatcc a 51

<210> 6162
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (6161 is other entry)

<221> misc_feature

<222> (0)...(0)
<223> Accession number cg44915044

<400> 6162
tgctcaaggg cgcggttttc ttccttacgg acttcgcgcg caagctatcc a 51

<210> 6163
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (6164 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44915044

<400> 6163
acttcgcgcg caagctatcc atcccctccg agctggagtt tatggccgtg t 51

<210> 6164
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (6163 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44915044

<400> 6164
acttcgcgcg caagctatcc atcccctccg agctggagtt tatggccgtg t 51

<210> 6165
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (6166 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44915044

<400> 6165
gcgcaagcta tccatcccct ccgagctgga gtttatggcc gtgtcctctt a 51

<210> 6166
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (6165 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44915044

<400> 6166
gcgcaagcta tccatccct ccgagttgga gtttatggcc gtgtcctctt a 51

<210> 6167
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (6168 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44915044

<400> 6167
agctggagtt tatggccgtg tcctcttatg gcgcctccac ttcttcttcc g 51

<210> 6168
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (6167 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44915044

<400> 6168
agctggagtt tatggccgtg tcctcgatg gcgcctccac ttcttcttcc g 51

<210> 6169
<211> 51
<212> DNA
<213> Homo sapiens

<220>

<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (6170 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44915044

<400> 6169
atggcgccctc cactttcttct tccggcgtgg tgcgcatttt gaaggacttg g 51

<210> 6170
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (6169 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44915044

<400> 6170
atggcgccctc cactttcttct tccggcgtgg tgcgcatttt gaaggacttg g 51

<210> 6171
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (6172 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44915044

<400> 6171
caatcctcat tggagaagag gaactacaaa accgcgtcca agagctcgcg g 51

<210> 6172
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (6171 is other entry)

<221> misc_feature
<222> (0)...(0)

<223> Accession number cg44915044

<400> 6172

caatcctcat tggagaagag gaactgcaaa accgcgtcca agagctcgcg g 51

<210> 6173

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (6174 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44915062

<400> 6173

gtctccaata tgtaagatag cacagcttat tcagaaaata acgagattga g 51

<210> 6174

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (6173 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44915062

<400> 6174

gtctccaata tgtaagatag cacagtttat tcagaaaata acgagattga g 51

<210> 6175

<211> 46

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (21)...(0)

<223> 1 of 2 allelic variants (6176 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44915070

<400> 6175

ntccggagcg gttcagcgcg aatgccgctt tctacgcgtg agagcg 46

<210> 6176

<211> 46
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (21)...(0)
<223> 2 of 2 allelic variants (6175 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44915070

<400> 6176
ntccggagcg gttcagcgcg gatgccgctt tctacgcgtg agagcg

46

<210> 6177
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (6178 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44915080

<400> 6177
tgtgtccaca tgtgaccgga gtgcctgcgc ccagccgccc gtgacacttc c

51

<210> 6178
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (6177 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44915080

<400> 6178
tgtgtccaca tgtgaccgga gtgcccgcgc ccagccgccc gtgacacttc c

51

<210> 6179
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature

<222> (26)...(0)
<223> 1 of 2 allelic variants (6180 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44915080

<400> 6179
cggagtgcct gcgcccagcc gcccgtagaca cttccgggct gagacgcggt c 51

<210> 6180
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (6179 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44915080

<400> 6180
cggagtgcct gcgcccagcc gcccgtagaca cttccgggct gagacgcggt c 51

<210> 6181
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (6182 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44915080

<400> 6181
gcccgtagaca cttccgggct gagacgcggt ctggtcttaa ggggcccgtc a 51

<210> 6182
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (6181 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44915080

<400> 6182
gccccgtgaca cttccgggct gagacacggg ctggtcttaa ggggcccgtc a 51

<210> 6183
<211> 50
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (6184 is other entry)

<221> misc_feature
<222> (25)...(26)
<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44915080

<400> 6183
cgggctgaga cgcggtctgg tcttaagggg cccgtcacag cgatggggccc 50

<210> 6184
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (6183 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44915080

<400> 6184
cgggctgaga cgcggtctgg tcttacaggg gcccgtcaca gcgatggggc c 51

<210> 6185
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (6186 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44915149

<400> 6185

tatcgatccc cggttccgga ccctccacga tcatcaaadc caccagaaaa a 51

<210> 6186

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (6185 is other entry)

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<222> (0)...(0)

<223> Accession number cg44915149

<400> 6186

tatcgatccc cggttccgga ccctcgacga tcatcaaadc caccagaaaa a 51

<210> 6187

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (6188 is other entry)

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<222> (0)...(0)

<223> Accession number cg44915149

<400> 6187

tactgatgcc gatcgacagg ccgagtgaga gctgggtgcc gctgtgacca a 51

<210> 6188

<211> 51

<212> DNA

<213> Homo sapiens

<220>

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<223> 2 of 2 allelic variants (6187 is other entry)

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<222> (0)...(0)

<223> Accession number cg44915149

<400> 6188

tactgatgcc gatcgacagg ccgagcgaga gctgggtgcc gctgtgacca a 51

<210> 6189

<211> 51

<212> DNA

<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (6190 is other entry)

<221> misc_feature
<222> (0)...(0)
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<400> 6189
cttgaacgtc agtcggtgcc cggtaggcgt actttcccag acggccgtca c 51

<210> 6190
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (6189 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44915354

<400> 6190
cttgaacgtc agtcggtgcc cggtaacggt actttcccag acggccgtca c 51

<210> 6191
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (6192 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44915354

<400> 6191
cggtgcccg taggcgtact ttcccagacg gccgtcacct cgaatggcgt c 51

<210> 6192
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (6191 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44915354

<400> 6192
cgggtgcccgg taggcgtact ttccccgacg gccgtcacct cgaatggcgt c 51

<210> 6193
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (6194 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44915354

<400> 6193
ccggtaggcg tactttccca gacggccgtc acctcgaatg gcgtcaaadc g 51

<210> 6194
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (6193 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44915354

<400> 6194
ccggtaggcg tactttccca gacggtcgtc acctcgaatg gcgtcaaadc g 51

<210> 6195
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (6196 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44915354

<400> 6195
gacccgcagt tacccttctt catcaaatgg acctctccgg ccgaactgtt g 51

<210> 6196
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (6195 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44915354

<400> 6196
gaccgcgagt tacccttctt catcacatgg acctctccgg ccgaactggt g 51

<210> 6197
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (6198 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44915354

<400> 6197
agttaccctt cttcatcaaa tggacctctc cggccgaact gttgccttcc a 51

<210> 6198
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (6197 is other entry)

<221> misc_feature
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<223> Accession number cg44915354

<400> 6198
agttaccctt cttcatcaaa tggacttctc cggccgaact gttgccttcc a 51

<210> 6199
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (6200 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44915354

<400> 6199
cttcttcacac aaatggacac ctccggccga actgttgccct tccagcctga g 51

<210> 6200
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (6199 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44915354

<400> 6200
cttcttcacac aaatggacac ctccgaccga actgttgccct tccagcctga g 51

<210> 6201
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (6202 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44915354

<400> 6201
ccgaactggt gccttcacac ctgagtggca acgtcacctt ggcctccgac g 51

<210> 6202
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (6201 is other entry)

<221> misc_feature

<222> (0)...(0)
<223> Accession number cg44915354

<400> 6202
ccgaactgtt gccttcacgc ctgagcggca acgtcacctt ggcctccgtc g 51

<210> 6203
<211> 49
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (6204 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44915354

<400> 6203
ttccagcctg agtggcaacg tcaccttggc ctccgtcgaa atgtccgga 49

<210> 6204
<211> 49
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (6203 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44915354

<400> 6204
ttccagcctg agtggcaacg tcaccttggc ctccgtcgaa atgtccgga 49

<210> 6205
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (6206 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44915354

<400> 6205
gtgcacgcca agagcagggc ggcggttggg tgggatgggc cgtctccgtc g 51

<210> 6206
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (6205 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44915354

<400> 6206
gtgcacgccca agagcagggc ggcggctgga tgggatgggc cgtctccgtc g 51

<210> 6207
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (6208 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44915354

<400> 6207
aagagcaggg cggcggttgg atgggatggg ccgtctccgt cgacgatctc g 51

<210> 6208
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (6207 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44915354

<400> 6208
aagagcaggg cggcggttgg atggggtggg ccgtctccgt cgacgatctc g 51

<210> 6209
<211> 51
<212> DNA
<213> Homo sapiens

<220>

<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (6210 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44915354

<400> 6209
gggccgtctc cgtcgacgat ctcgcccctt atgagaagcg tcttgaacgt c 51

<210> 6210
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (6209 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44915354

<400> 6210
gggccgtctc cgtcgacgat ctcgctcctt atgagaagcg tcttgaacgt c 51

<210> 6211
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (6212 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44915354

<400> 6211
acgatctcgc cccttatgag aagcgtcttg aacgtcagtc ggtgcccggt a 51

<210> 6212
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (6211 is other entry)

<221> misc_feature
<222> (0)...(0)

<223> Accession number cg44915354

<400> 6212

acgatctcgc cccttatgag aagcgcttg aacgtcagtc ggtgcccggt a

51

<210> 6213

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (6214 is other entry)

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<222> (0)...(0)

<223> Accession number cg44915354

<400> 6213

tcgcccctta tgagaagcgt cttgaacgtc agtcggtgcc cggtaggcgt a

51

<210> 6214

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (6213 is other entry)

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<222> (0)...(0)

<223> Accession number cg44915354

<400> 6214

tcgcccctta tgagaagcgt cttgagcgtc agtcggtgcc cggtaggcgt a

51

<210> 6215

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (6216 is other entry)

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<222> (0)...(0)

<223> Accession number cg44915354

<400> 6215

ccccttatga gaagcgtctt gaacgtcagt cggtgcccgg taggcgtact t

51

<210> 6216

<211> 51
<212> DNA
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<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (6215 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44915354

<400> 6216
ccccttatga gaagcgtctt gaacgccagt cggtgcccgg taggcgtact t 51

<210> 6217
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (6218 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44915425

<400> 6217
tccaaccgat gatcaggaat gggacttgct gttgacggta accaaatggt g 51

<210> 6218
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (6217 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44915425

<400> 6218
tccaaccgat gatcaggaat gggacctgct gttgacggta accaaatggt g 51

<210> 6219
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (6220 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44915425

<400> 6219
cgcgatggtg cgccatagtc acgaagcagc tggcccctca gccagggaaa a 51

<210> 6220
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (6219 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44915425

<400> 6220
cgcgatggtg cgccatagtc acgaaacagc tggcccctca gccagggaaa a 51

<210> 6221
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (6222 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44915425

<400> 6221
gtgcgccata gtcacgaagc agctggcccc tcagccaggg aaaagcatcc g 51

<210> 6222
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (6221 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44915425

<400> 6222
gtgcgccata gtcacgaagc agctgtcccc tcagccaggg aaaagcatcc g 51

<210> 6223
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (6224 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44916342

<400> 6223
gatgatgagg aaaaatgtgg ttgtcgtatt atgtgtccgg tcaggcctct g 51

<210> 6224
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (6223 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44916342

<400> 6224
gatgatgagg aaaaatgtgg ttgtcatatt atgtgtccgg tcaggcctct g 51

<210> 6225
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (6226 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44916489

<400> 6225
cggcgccgcg ctccttgatg gaagccgcct tcttgtcgtc gacaaacccc a 51

<210> 6226
<211> 51

<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (6225 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44916489

<400> 6226
cggcgccgc ctccttgatg gaagcggcct tcttgctgc gacaaacccc a

51

<210> 6227
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (6228 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44916489

<400> 6227
gcctccttga tggaagccgc cttcttgctg tcgacaaacc ccaaacagac a

51

<210> 6228
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (6227 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44916489

<400> 6228
gcctccttga tggaagccgc cttctcgctg tcgacaaacc ccaaacagac a

51

<210> 6229
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)

<223> 1 of 2 allelic variants (6230 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44917150

<400> 6229

ggtgcccccg gcatgaggtg cggcctccag ggcgcctcct gccggccccg a

51

<210> 6230

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (6229 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44917150

<400> 6230

ggtgcccccg gcatgaggtg cggccccag ggcgcctcct gccggccccg a

51

<210> 6231

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (6232 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44917278

<400> 6231

agctttgggg gagggtaatt cctgccagga agctgggtcc accagtcctg c

51

<210> 6232

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (6231 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44917278

<400> 6232
agctttgggg gagggtaatt cctgctagga agctgggtcc accagtcctg c 51

<210> 6233
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (6234 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44917278

<400> 6233
tgggggaggg taattcctgc caggaagctg gttccaccag tcctgctgta g 51

<210> 6234
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (6233 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44917278

<400> 6234
tgggggaggg taattcctgc caggaggctg gttccaccag tcctgctgta g 51

<210> 6235
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (6236 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44917461

<400> 6235
ttggaagatc tgaagctcccg ccacgcctt cttggcttca ccggcgtcaa c 51

<210> 6236
<211> 51
<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (6235 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44917461

<400> 6236

ttggaagatc tgagctcccg ccatcacctt cttggcttca ccggcgtaa c

51

<210> 6237

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (6238 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44917712

<400> 6237

atcgccatc tgcaccccca tgggggctag agccctttac ttctccttc a

51

<210> 6238

<211> 50

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (6237 is other entry)

<221> misc_feature

<222> (25)...(26)

<223> Nucleotide deleted between bases 25 and 26

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44917712

<400> 6238

atcgccatc tgcaccccca tgggggctaga gccctttact tcctccttca

50

<210> 6239

<211> 51

<212> DNA

<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (6240 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44917712

<400> 6239
agtagacaat aaataactgc ttttccggca gaggctaggt tgcctgggg g 51

<210> 6240
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (6239 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44917712

<400> 6240
agtagacaat aaataactgc ttttctggca gaggctaggt tgcctgggg g 51

<210> 6241
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (6242 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44918306

<400> 6241
atcttttggt ttgctttttt ttttttaaaa aagggtcccag gaatatacag c 51

<210> 6242
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<400> 6242
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<210> 6243
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<223> 1 of 2 allelic variants (6244 is other entry)

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tcggtctggg aagcgggtgac gtaggactcg tcggcagacg aggtgcccac g 51

<210> 6244
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<210> 6245
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<221> misc_feature
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<400> 6245
tatctcttca acagatccgg gtggccattg accatctacg cagcaacggg g 51

<210> 6246
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<223> 2 of 2 allelic variants (6245 is other entry)

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<400> 6246
tatctcttca acagatccgg gtggctattg accatctacg cagcaacggg g 51

<210> 6247
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<221> misc_feature
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gaagtcagcg cctgtactct ttccgtgaca tcctcatgct caaggttgct a 51

<210> 6248
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<223> 2 of 2 allelic variants (6247 is other entry)

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<223> Accession number cg44918799

<400> 6248
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<210> 6249
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<212> DNA
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<223> 1 of 2 allelic variants (6250 is other entry)

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<210> 6250
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<210> 6251
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<210> 6252
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<223> 2 of 2 allelic variants (6251 is other entry)

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tcatgctcaa ggttggtcaag cgtctccttg acgctgggggt atctcttcaa c 51

<210> 6253
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<212> DNA
<213> Homo sapiens

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<223> 1 of 2 allelic variants (6254 is other entry)

<221> misc_feature
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<223> Accession number cg44918827

<400> 6253
tcagcaagaa caagacgcct gccctcaagg gctcgccgca gcgtcgtggc g 51

<210> 6254
<211> 51
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<223> 2 of 2 allelic variants (6253 is other entry)

<221> misc_feature
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<400> 6254
tcagcaagaa caagacgcct gccctgaagg gctcgccgca gcgtcgtggc g 51

<210> 6255
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accgctcgag ttgctgagag aaggtctgcc cgtgcggagt tcgcacggca c 51

<210> 6256
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 2 of 2 allelic variants (6255 is other entry)

<221> misc_feature
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<223> Accession number cg44918827

<400> 6256
accgctcgag ttgctgagag aaggtttgcc cgtgcggagt tcgcacggca c 51

<210> 6257
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<212> DNA
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<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (6258 is other entry)

<221> misc_feature
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<223> Accession number cg44918840

<400> 6257
tagctgaggt gaggagcagg gacggaatcg ggcagcagctc gcgagtggcg t 51

<210> 6258
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (6257 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44918840

<400> 6258
tagctgaggt gaggagcagg gacgggatcg ggcagcagctc gcgagtggcg t 51

<210> 6259
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<213> Homo sapiens

<220>

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<221> misc_feature

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<223> Accession number cg44918840

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cgcgagtggc gtggcgggtcc gagagtacaa tgatggtagt tccctcagca a

51

<210> 6260

<211> 51

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<223> 2 of 2 allelic variants (6259 is other entry)

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<223> Accession number cg44918840

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51

<210> 6261

<211> 51

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<222> (26)...(0)

<223> 1 of 2 allelic variants (6262 is other entry)

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<213> Homo sapiens

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<223> 2 of 2 allelic variants (6261 is other entry)

<221> misc_feature
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<223> Accession number cg44918840

<400> 6262
gggtccgagag tacaatgatg gtagtcccct cagcaatggc ctcagaaacc t 51

<210> 6263
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<221> misc_feature
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<223> Accession number cg44918840

<400> 6263
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<210> 6264
<211> 51
<212> DNA
<213> Homo sapiens

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<222> (26)...(0)
<223> 2 of 2 allelic variants (6263 is other entry)

<221> misc_feature
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<223> Accession number cg44918840

<400> 6264
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<210> 6265
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<212> DNA
<213> Homo sapiens

<220>
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<221> misc_feature
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<400> 6265

gacgcaagcc ctcgccaccc tcttcaacaa caaagaggcc cctcaccacg t

51

<210> 6266

<211> 51

<212> DNA

<213> Homo sapiens

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<222> (26)...(0)

<223> 2 of 2 allelic variants (6265 is other entry)

<221> misc_feature

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<223> Accession number cg44918840

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gacgcaagcc ctcgccaccc tcttcgacaa caaagaggcc cctcaccacg t

51

<210> 6267

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 1 of 2 allelic variants (6268 is other entry)

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<222> (0)...(0)

<223> Accession number cg44918840

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gaacgtcacc agcctcgacg acgagaccgg cccgggtacg ggtcttctcc c

51

<210> 6268

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 2 of 2 allelic variants (6267 is other entry)

<221> misc_feature

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<223> Accession number cg44918840

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51

<210> 6269

<211> 51

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<213> Homo sapiens

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<223> 1 of 2 allelic variants (6270 is other entry)

<221> misc_feature
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<223> Accession number cg44920731

<400> 6269
tggaggaaaa tatgtacatc aatgcgcacc agtgatcaga aaacccccag g 51

<210> 6270
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 2 of 2 allelic variants (6269 is other entry)

<221> misc_feature
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tggaggaaaa tatgtacatc aatgcacacc agtgatcaga aaacccccag g 51

<210> 6271
<211> 51
<212> DNA
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<223> 1 of 2 allelic variants (6272 is other entry)

<221> misc_feature
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<223> Accession number cg44920731

<400> 6271
aaatatgtac atcaatgcgc accagtgatc agaaaacccc caggaaccga a 51

<210> 6272
<211> 51
<212> DNA
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<223> 2 of 2 allelic variants (6271 is other entry)

<221> misc_feature
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<400> 6272
aaatatgtac atcaatgcgc accagggatc agaaaacccc caggaaccca a 51

<210> 6273
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 1 of 2 allelic variants (6274 is other entry)

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<400> 6273
tacatcaatg cgcaccagtg atcagaaaac cccaggaac ccaagcaagt g 51

<210> 6274
<211> 50
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<400> 6274
tacatcaatg cgcaccagtg atcagaaacc cccaggaacc caagcaagtg 50

<210> 6275
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 1 of 2 allelic variants (6276 is other entry)

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<223> Accession number cg44920731

<400> 6275

gcaccagtga tcagaaaacc cccaggaacc caagcaagtg ggaactgagg g

51

<210> 6276

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 2 of 2 allelic variants (6275 is other entry)

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<223> Accession number cg44920731

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<210> 6277

<211> 51

<212> DNA

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<223> 1 of 2 allelic variants (6278 is other entry)

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51

<210> 6278

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 2 of 2 allelic variants (6277 is other entry)

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<223> Accession number cg44920731

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gatcagaaaa cccccaggaa cccaaccaag tgggaactga gggggccggc t

51

<210> 6279

<211> 51
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<400> 6279
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<210> 6280
<211> 50
<212> DNA
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<223> 2 of 2 allelic variants (6279 is other entry)

<221> misc_feature
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<400> 6280
ggctcctcat cagctgggga aaaggaaaat gggcctcaca gaagccataa 50

<210> 6281
<211> 51
<212> DNA
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<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (6282 is other entry)

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<400> 6281
cctcatcagc tggggaaaag ggaaaatggg cctcacagaa gccataacag g 51

<210> 6282
<211> 51
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<213> Homo sapiens

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<223> 2 of 2 allelic variants (6281 is other entry)

<221> misc_feature

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cctcatcagc tggggaaaag ggaaagtggg cctcacagaa gccataacag g

51

<210> 6283

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 1 of 2 allelic variants (6284 is other entry)

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<223> Accession number cg44920731

<400> 6283

aatgggcct cacagaagcc ataacagggt ggaaagagcg aggctgcagt c

51

<210> 6284

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 2 of 2 allelic variants (6283 is other entry)

<221> misc_feature

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<223> Accession number cg44920731

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51

<210> 6285

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (6286 is other entry)

<221> misc_feature
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<400> 6285
tcacagaagc cataacaggg tggaaagagc gaggtgcag tccacagggg t 51

<210> 6286
<211> 51
<212> DNA
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<221> misc_feature
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<400> 6286
tcacagaagc cataacaggg tggaaaggagc gaggtgcag tccacagggg t 51

<210> 6287
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (6288 is other entry)

<221> misc_feature
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<400> 6287
gaagccataa caggggtggaa agagcgaggc tgcagtccac aggggttggtg t 51

<210> 6288
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (6287 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44920731

<400> 6288

gaagccataa caggggtggaa agagcaaggc tgcagtccac aggggttggtg t 51

<210> 6289

<211> 51

<212> DNA

<213> Homo sapiens

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<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (6290 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44920731

<400> 6289

agccataaca ggggtggaaag agcgaggctg cagtccacag gggttgtgtg a 51

<210> 6290

<211> 51

<212> DNA

<213> Homo sapiens

<220>

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<222> (26)...(0)

<223> 2 of 2 allelic variants (6289 is other entry)

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<222> (0)...(0)

<223> Accession number cg44920731

<400> 6290

agccataaca ggggtggaaag agcgaagctg cagtccacag gggttgtgtg a 51

<210> 6291

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (6292 is other entry)

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<210> 6292

<211> 50

<212> DNA

<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (6291 is other entry)

<221> misc_feature
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<221> misc_feature
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<223> Accession number cg44920731

<400> 6292
gttgtgttga gtggcttgta tttctctctg cagggggagt ggcattctct 50

<210> 6293
<211> 51
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<220>
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<223> 1 of 2 allelic variants (6294 is other entry)

<221> misc_feature
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<223> Accession number cg44920877

<400> 6293
gggtctatgc cccgcggcgc ttggcctcgc ccacaccccc ggcacctgc c 51

<210> 6294
<211> 50
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (6293 is other entry)

<221> misc_feature
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<221> misc_feature
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<400> 6294
gggtctatgc cccgcggcgc ttggctcgc cacacccccg ggcacctgc 50

<210> 6295
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<212> DNA
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<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (6296 is other entry)

<221> misc_feature
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<400> 6295
ggccgttggc ctgcgccaca cccccggccc cactgcgggt ggagagacgt c

51

<210> 6296
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (6295 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg44920877

<400> 6296
ggccgttggc ctgcgccaca cccccggccc cactgcgggt ggagagacgt c

51

<210> 6297
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 1 of 2 allelic variants (6298 is other entry)

<221> misc_feature
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<223> Accession number cg43959936

<400> 6297
tgtgtgatgg ggacatggag aagcccatcc aggtcatgtg ctacgactat g

51

<210> 6298
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> Accession number cg43959936

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51

<210> 6299

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<223> Accession number cg34407554

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51

<210> 6300

<211> 51

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<223> Accession number cg34407554

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51

<210> 6301

<211> 51

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<223> 1 of 2 allelic variants (6302 is other entry)

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<223> Accession number cg34407554

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acgtcgctgg caaacatcgc gactgcgcg c aagtaggcatt tgaatcgatt c 51

<210> 6304
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<212> DNA
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<223> 2 of 2 allelic variants (6303 is other entry)

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acgtcgctgg caaacatcgc gactgtgcgc aagtaggcatt tgaatcgatt c 51

<210> 6305
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<223> 1 of 2 allelic variants (6306 is other entry)

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51

<210> 6306

<211> 51

<212> DNA

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<210> 6307

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51

<210> 6308

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<210> 6309
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<210> 6310
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<210> 6311
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<210> 6312

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<223> Accession number cg29344427

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<210> 6313

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<223> 1 of 2 allelic variants (6314 is other entry)

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<222> (0)...(0)

<223> Accession number cg42716656

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51

<210> 6314

<211> 51

<212> DNA

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<210> 6315

<211> 51

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<210> 6316
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<210> 6317
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gaggctggcc tgtggtgatg tcgtagcaat gaccttcag taacaggtat t 51

<210> 6318
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gaggctggcc tgtggtgatg tcgtaacaat gaccttccag taacaggtat t 51

<210> 6319
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<210> 6320
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<210> 6321
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<400> 6321
cgtagcaatg accttccagt aacaggtatt ccagctcata ctcagcagcc a 51

<210> 6322
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<400> 6322
cgtagcaatg accttccagt aacagatatt ccagctcata ctcagcagcc a 51

<210> 6323
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<400> 6323
gccactacac tgtccacctc ttctaaataa atattatcaa gatcatatgg t 51

<210> 6324
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<210> 6325
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ccacctcttc taaataaata ttatcaagat catatgggtgt tctgacagat t 51

<210> 6326
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ccacctcttc taaataaata ttatccagat catatgggtgt tctgacagat t 51

<210> 6327
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<400> 6327
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<210> 6328
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gtgttctgac agattctacc atccaactct caggtgtggt caaattcaaa g 51

<210> 6329
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<210> 6330
<211> 51
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accatccagc tctcaggtgt gttcagattc aaagtaaaca gcggagactg a 51

<210> 6331
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<400> 6331
aacagcggag actgaggcat atccaaaaat tttgcgatcg gacccttagc a 51

<210> 6332

<211> 51

<212> DNA

<213> Homo sapiens

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51

<210> 6333

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 1 of 2 allelic variants (6334 is other entry)

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<223> Accession number cg42716656

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51

<210> 6334

<211> 51

<212> DNA

<213> Homo sapiens

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<222> (26)...(0)

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<221> misc_feature

<222> (0)...(0)

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51

<210> 6335

<211> 51

<212> DNA

<213> Homo sapiens

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<400> 6335
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<210> 6336
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<400> 6336
actgaggcat atccaaaaat ttgctatcg gacccttagc aaaactactg t 51

<210> 6337
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<400> 6337
atattgcgat cggaccctta gcaaaactac tgtctgaagt gaaagaaatc t 51

<210> 6338
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<223> Accession number cg42716656

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attttgcgat cggaccctta gcaaagctac tgtctgaagt gaaagaaatc t

51

<210> 6339

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<223> Accession number cg42716656

<400> 6339

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51

<210> 6340

<211> 51

<212> DNA

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<223> 2 of 2 allelic variants (6339 is other entry)

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51

<210> 6341

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<223> 1 of 2 allelic variants (6342 is other entry)

<221> misc_feature

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<223> Accession number cg27241188

<400> 6341

cggtgtcgag gtccgtgaag gccgggctcc agttccccgt cggccgcac g

51

<210> 6342

<211> 51
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<221> misc_feature
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<400> 6342
cgggtgtcgag gtccgtgaag gccgggtctcc agttccccgt cggccgcac g 51

<210> 6343
<211> 51
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<223> 1 of 2 allelic variants (6344 is other entry)

<221> misc_feature
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<223> Accession number cg20725546

<400> 6343
ttgccaaagga ggtagaggtc ctccaaggcc tgaccgccga tccgaagcgg c 51

<210> 6344
<211> 51
<212> DNA
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<223> 2 of 2 allelic variants (6343 is other entry)

<221> misc_feature
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<223> Accession number cg20725546

<400> 6344
ttgccaaagga ggtagaggtc ctccagggcc tgaccgccga tccgaagcgg c 51

<210> 6345
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<221> misc_feature
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<223> Accession number cg20725546

<400> 6345
gtatggcgta caccttcctc aaggcgaagg gtctcgaggt cggtgactcc c 51

<210> 6346
<211> 51
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<223> 2 of 2 allelic variants (6345 is other entry)

<221> misc_feature
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<223> Accession number cg20725546

<400> 6346
gtatggcgta caccttcctc aaggccaagg gtctcgaggt cggtgactcc c 51

<210> 6347
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<212> DNA
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<223> 1 of 2 allelic variants (6348 is other entry)

<221> misc_feature
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<223> Accession number cg44013488

<400> 6347
agaaggccaa agtccaggca tccacaggtc ctgcatctat agacttcac t 51

<210> 6348
<211> 51
<212> DNA
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<223> 2 of 2 allelic variants (6347 is other entry)

<221> misc_feature
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<400> 6348
agaaggccaa agtccaggca tccaccgggc ctgcatctat agacttcac t 51

<210> 6349
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<223> 1 of 2 allelic variants (6350 is other entry)

<221> misc_feature
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<400> 6349
agagacgaat atgagcatct agagagctgg aggcagcaat gggcaggggtg t 51

<210> 6350
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agagacgaat atgagcatct agagaactgg aggcagcaat gggcaggggtg t 51

<210> 6351
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<210> 6352
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<212> DNA
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ttgcaatattc aaaaatattg atgatcccat ctatggctcc actggctagg t 51

<210> 6353
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<223> 1 of 2 allelic variants (6354 is other entry)

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<400> 6353
tattgatgat tccatctatg gctccactgg ctaggtatctt cccatcagga c 51

<210> 6354
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tattgatgat tccatctatg gctccgctgg ctaggtatctt cccatcagga c 51

<210> 6355
<211> 51
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<223> 1 of 2 allelic variants (6356 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44013488

<400> 6355

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51

<210> 6356

<211> 51

<212> DNA

<213> Homo sapiens

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51

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<223> 1 of 2 allelic variants (6358 is other entry)

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<223> Accession number cg44013488

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51

<210> 6358

<211> 51

<212> DNA

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<223> Accession number cg44013488

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<210> 6360
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<210> 6361
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<223> Accession number cg42512386

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<210> 6362
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51

<210> 6363

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<223> Accession number cg29345567

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51

<210> 6364

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<223> Accession number cg29345567

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<223> 1 of 2 allelic variants (6366 is other entry)

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<210> 6366
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<210> 6367
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<223> Accession number cg29358731

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<210> 6368
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51

<210> 6369

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<223> Accession number cg29358731

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51

<210> 6370

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51

<210> 6371

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51

<210> 6372

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<400> 6372
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<210> 6373
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<223> 1 of 2 allelic variants (6374 is other entry)

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<210> 6374
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<400> 6374
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<210> 6375
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<223> 1 of 2 allelic variants (6376 is other entry)

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<210> 6376
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<400> 6376
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<210> 6377
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<210> 6378
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<210> 6379
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<210> 6380
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<210> 6381
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ggttgagaaa ctcaggaaag tcgatgggtc cattgccatc agcatcaacc t 51

<210> 6382
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<400> 6382
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<210> 6383
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<210> 6384
<211> 48
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<400> 6384
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<210> 6385
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<210> 6386
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<210> 6387
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<223> Accession number cg43976004

<400> 6387
aggacgacga cgacacgcgt gaaggggagg aggagctaga ggacgacgac g 51

<210> 6388
<211> 51
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<223> Accession number cg43976004

<400> 6388
aggacgacga cgacacgcgt gaaggagagg aggagctaga ggacgacgac g 51

<210> 6389
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<223> Accession number cg44127439

<400> 6389
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<210> 6390
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<223> Accession number cg44127439

<400> 6390
cggcgcgccgc ctccttgatg gaagcgcgcct tcttgtcgtc gacaaacccc a 51

<210> 6391
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<400> 6391
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<210> 6392
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<400> 6392
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<210> 6393
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<400> 6393
agagagtcca cactggagag aaacctata gatgttggtg atgtgggaag g 51

<210> 6394
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<400> 6394
agagagtcca cactggagag aaacctata gatgttggtg atgtgggaag g 51

<210> 6395
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<223> Accession number cg32152874

<400> 6395

gcggcaaggt gtacgtgtcc atgccggcca tggccatgca cctgctcacg c

51

<210> 6396

<211> 51

<212> DNA

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<223> 2 of 2 allelic variants (6395 is other entry)

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<223> Accession number cg32152874

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<210> 6397

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<223> Accession number cg32152874

<400> 6397

acgacctgcg ccacaagtgc ggcgtgtgcg gcaaagcctt ctgcgggccc t

51

<210> 6398

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<212> DNA

<213> Homo sapiens

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<223> 2 of 2 allelic variants (6397 is other entry)

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<223> Accession number cg32152874

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51

<210> 6399

<211> 51
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gggacgcgtg cggcgagtg gcgcaaacat acgccacgtc gtcgaacctg a 51

<210> 6400
<211> 51
<212> DNA
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<223> 2 of 2 allelic variants (6399 is other entry)

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gggacgcgtg cggcgagtg gcgaagacat acgccacgtc gtcgaacctg a 51

<210> 6401
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<212> DNA
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<223> 1 of 2 allelic variants (6402 is other entry)

<221> misc_feature
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<400> 6401
ttccataacg gctggaagct gcaaagaagc ttgtgatttg gtaaccaaag c 51

<210> 6402
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 2 of 2 allelic variants (6401 is other entry)

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ttccataacg gctggaagct gcaaaaaagc ttgtgatttg gtaaccaaag c 51

<210> 6403
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<212> DNA
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<223> 1 of 2 allelic variants (6404 is other entry)

<221> misc_feature
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<223> Accession number cg43103077

<400> 6403
ctgcaaagaa gcttgtgatt tggtaaccaa agctggcata gtaagcatgc t 51

<210> 6404
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<212> DNA
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<223> 2 of 2 allelic variants (6403 is other entry)

<221> misc_feature
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<400> 6404
ctgcaaagaa gcttgtgatt tggtaaccaa agctggcata gtaagcatgc t 51

<210> 6405
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<223> 1 of 2 allelic variants (6406 is other entry)

<221> misc_feature
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<400> 6405
aaccaaagct ggcataagtaa gcatgctcca tgattgccat caactgaatg c 51

<210> 6406
<211> 51
<212> DNA
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<400> 6406
aaccaaagct ggcataagtaa gcatgttcca tgattgccat caactgaatg c 51

<210> 6407
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<212> DNA
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<400> 6407
tggcatagta agcatgctcc atgattgccca tcaactgaat gcagttgtat c 51

<210> 6408
<211> 51
<212> DNA
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tggcatagta agcatgctcc atgattgccca tcaactgaat gcagttgtat c 51

<210> 6409
<211> 51

<212> DNA
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<223> 1 of 2 allelic variants (6410 is other entry)

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<400> 6409
ttgccatcaa ctgaatgcag ttgtatccaa ggcctttgat tcttggtagt a 51

<210> 6410
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<400> 6410
ttgccatcaa ctgaatgcag ttgtacccaa ggcctttgat tcttggtagt a 51

<210> 6411
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<223> 1 of 2 allelic variants (6412 is other entry)

<221> misc_feature
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<223> Accession number cg20749088

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taggggtggc agtgccaatg tccatccgc gatacaccac catcgagtca g 51

<210> 6412
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<212> DNA
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<223> 2 of 2 allelic variants (6411 is other entry)

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<223> Accession number cg20749088

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51

<210> 6413

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 1 of 2 allelic variants (6414 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg38434693

<400> 6413

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51

<210> 6414

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51

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<223> Accession number cg38434693

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<210> 6418
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51

<210> 6420

<211> 51

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<223> 2 of 2 allelic variants (6419 is other entry)

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<223> Accession number cg38434693

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51

<210> 6421

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<223> Accession number cg38434693

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51

<210> 6422

<211> 51

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<223> 2 of 2 allelic variants (6421 is other entry)

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<210> 6423
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<210> 6424
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<210> 6425
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<223> 1 of 2 allelic variants (6426 is other entry)

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gctacctgcc acaggatccc cgcgacccag acatggaaat gatcgcgagg g 51

<210> 6426

<211> 51

<212> DNA

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<210> 6427

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<223> 1 of 2 allelic variants (6428 is other entry)

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<222> (0)...(0)

<223> Accession number cg43919179

<400> 6427

taaacctccc gttgggagtg atcatataga tactgggagg tttgaaagga a 51

<210> 6428

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 2 of 2 allelic variants (6427 is other entry)

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<223> Accession number cg43919179

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taaacctccc gttgggagtg atcatgtaga tactgggagg tttgaaagga a 51

<210> 6429

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<223> Accession number cg25254092

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<210> 6430
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<400> 6430
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<210> 6431
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<223> 1 of 2 allelic variants (6432 is other entry)

<221> misc_feature
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<210> 6432
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<223> 2 of 2 allelic variants (6431 is other entry)

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<210> 6433
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<223> 1 of 2 allelic variants (6434 is other entry)

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<210> 6435
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<223> 1 of 2 allelic variants (6436 is other entry)

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tgcatttcca gtatgttgca agattcagag gagggtgactg aatgccattt g 51

<210> 6436
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<223> 2 of 2 allelic variants (6435 is other entry)

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tgcatttcca gtatgttgca agatttagag gaggggactg aatgccattt g 51

<210> 6437
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<223> 1 of 2 allelic variants (6438 is other entry)

<221> misc_feature
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<223> Accession number cg39517880

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<210> 6438
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<400> 6438
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<210> 6439
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<223> 1 of 2 allelic variants (6440 is other entry)

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<210> 6440
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<210> 6441
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<223> 1 of 2 allelic variants (6442 is other entry)

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cagtgggctg gatgacgtca tgaacatcgt cacgaggtga catcagcagg g 51

<210> 6442
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<223> 2 of 2 allelic variants (6441 is other entry)

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<210> 6443
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<223> 1 of 2 allelic variants (6444 is other entry)

<221> misc_feature
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<223> Accession number cg39517880

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tgacgtcatg aacatcgtca cgaggtgaca tcagcagggt gtagatggaa t 51

<210> 6444
<211> 51
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<223> 2 of 2 allelic variants (6443 is other entry)

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<223> Accession number cg39517880

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<210> 6445
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<221> misc_feature
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<223> Accession number cg29966014

<400> 6445
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<210> 6446
<211> 51
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gacgcaatca cgaactcctc gcgcagcgcc ggcgacacct cggggatggt c 51

<210> 6447
<211> 51
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<223> 1 of 2 allelic variants (6448 is other entry)

<221> misc_feature
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<223> Accession number cg43945782

<400> 6447
atgtggcttc cagatcctct gtcttggtgc ggagatgttc caagttttcc c 51

<210> 6448
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<223> 2 of 2 allelic variants (6447 is other entry)

<221> misc_feature
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<223> Accession number cg43945782

<400> 6448
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<210> 6449
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<223> 1 of 2 allelic variants (6450 is other entry)

<221> misc_feature
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<223> Accession number cg27837466

<400> 6449
gggtgatagc caccaactgc tcgcatcctt tgacgacctc gtcggtcaac c 51

<210> 6450
<211> 51
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<223> 2 of 2 allelic variants (6449 is other entry)

<221> misc_feature
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gggtgatagc caccaactgc tcgcaccctt tgacgacctc gtcggtcaac c 51

<210> 6451
<211> 51
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<223> 1 of 2 allelic variants (6452 is other entry)

<221> misc_feature
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catcctttga cgacctcgtc ggtcaaccgc ggaccttcgc atccaaagac c 51

<210> 6452
<211> 51
<212> DNA
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<223> 2 of 2 allelic variants (6451 is other entry)

<221> misc_feature
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<223> Accession number cg27837466

<400> 6452

catcctttga cgacctcgtc ggtagcccg ggaccttcg atccaaagac c

51

<210> 6453

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 1 of 2 allelic variants (6454 is other entry)

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<222> (0)...(0)

<223> Accession number cg43287601

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51

<210> 6454

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 2 of 2 allelic variants (6453 is other entry)

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acagggtttc accgtgttgg ccaggatggt cttgaactcc tgacctcaag t

51

<210> 6455

<211> 51

<212> DNA

<213> Homo sapiens

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<222> (26)...(0)

<223> 1 of 2 allelic variants (6456 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg39527289

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acggcgatgc caacttggct aaaggcgcca ccgtcggaat cttgcttcag g

51

<210> 6456

<211> 51
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<223> 2 of 2 allelic variants (6455 is other entry)

<221> misc_feature
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<223> Accession number cg39527289

<400> 6456
acggcgatgc caacttggt aaaggtgcca ccgtcggaat cttgcttcag g 51

<210> 6457
<211> 51
<212> DNA
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<223> 1 of 2 allelic variants (6458 is other entry)

<221> misc_feature
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<223> Accession number cg39527289

<400> 6457
aagtctccgc cgagatggcc aaccctgacg ccgactttga cgccctgatg g 51

<210> 6458
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 2 of 2 allelic variants (6457 is other entry)

<221> misc_feature
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<400> 6458
aagtctccgc cgagatggcc aaccccgacg ccgactttga cgccctgatg g 51

<210> 6459
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 1 of 2 allelic variants (6460 is other entry)

<221> misc_feature
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<223> Accession number cg39527289

<400> 6459
acgccgactt tgacgccctg atggcggaga tgggtgagct gcagaccgag c 51

<210> 6460
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 2 of 2 allelic variants (6459 is other entry)

<221> misc_feature
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acgccgactt tgacgccctg atggccgaga tgggtgagct gcagaccgag c 51

<210> 6461
<211> 51
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<213> Homo sapiens

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<223> 1 of 2 allelic variants (6462 is other entry)

<221> misc_feature
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aggctggcat ggagcgggaa ctggagaaca tcatccagga gacagagaaa g 51

<210> 6462
<211> 51
<212> DNA
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<223> 2 of 2 allelic variants (6461 is other entry)

<221> misc_feature
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aggctggcat ggagcgggaa ctggaaaaca tcatccagga gacagagaaa g 51

<210> 6463
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<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (6464 is other entry)

<221> misc_feature
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<223> Accession number cg43294390

<400> 6463
caccataata atctggtgca tcattagggt ctactggttc aaggaaaggc c 51

<210> 6464
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<223> 2 of 2 allelic variants (6463 is other entry)

<221> misc_feature
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caccataata atctggtgca tcattggggt ctactggttc aaggaaaggc c 51

<210> 6465
<211> 51
<212> DNA
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<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (6466 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg34394308

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<210> 6466
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<212> DNA
<213> Homo sapiens

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<223> 2 of 2 allelic variants (6465 is other entry)

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<223> Accession number cg34394308

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<210> 6467
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<223> 1 of 2 allelic variants (6468 is other entry)

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tcatccagca cggctctgacg acaaacgga aatcaatgga gtggtttaag c 51

<210> 6468
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<221> misc_feature
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<223> Accession number cg34394308

<400> 6468
tcatccagca cggctctgacg acaaaccgga aatcaatgga gtggtttaag c 51

<210> 6469
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<223> 1 of 2 allelic variants (6470 is other entry)

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<223> Accession number cg34394308

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51

<210> 6470

<211> 51

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<213> Homo sapiens

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<223> 2 of 2 allelic variants (6469 is other entry)

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<223> Accession number cg34394308

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51

<210> 6471

<211> 51

<212> DNA

<213> Homo sapiens

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51

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<223> Accession number cg43129081

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<210> 6474
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<223> 1 of 2 allelic variants (6476 is other entry)

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<223> Accession number cg42868441

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<223> 2 of 2 allelic variants (6475 is other entry)

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51

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<223> 1 of 2 allelic variants (6480 is other entry)

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<210> 6480
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<210> 6481
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<210> 6482
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<210> 6483

<211> 51

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<210> 6484

<211> 51

<212> DNA

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<223> 2 of 2 allelic variants (6483 is other entry)

<221> misc_feature

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<223> Accession number cg29256713

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<210> 6485

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<210> 6486

<211> 51

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<210> 6487
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<210> 6488
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<210> 6489
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<223> Accession number cg33204850

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<210> 6490
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<223> Accession number cg41616497

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<210> 6492
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<223> 2 of 2 allelic variants (6491 is other entry)

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<223> 1 of 2 allelic variants (6494 is other entry)

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51

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<223> 2 of 2 allelic variants (6493 is other entry)

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51

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<223> 1 of 2 allelic variants (6496 is other entry)

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<223> Accession number cg30785452

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51

<210> 6496

<211> 51

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<210> 6498
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<223> 2 of 2 allelic variants (6497 is other entry)

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<400> 6498
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<210> 6499
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<210> 6500
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<210> 6501
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<223> 1 of 2 allelic variants (6502 is other entry)

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<210> 6502
<211> 51
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<223> Accession number cg35929441

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<210> 6503
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<400> 6503
gaagaaaatc taggcaataa cattcaggac ataggcatgg gcaaagattt c 51

<210> 6504
<211> 51
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<210> 6505
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<223> 1 of 2 allelic variants (6506 is other entry)

<221> misc_feature
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<223> Accession number cg42667624

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<210> 6506
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<400> 6506
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<210> 6507
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<223> Accession number cg34997905

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<210> 6508
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<210> 6509
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<223> 1 of 2 allelic variants (6510 is other entry)

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<223> Accession number cg43967284

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<210> 6510

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<221> misc_feature

<222> (0)...(0)

<223> Accession number cg29242150

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<210> 6512

<211> 51

<212> DNA

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<223> Accession number cg29242150

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<210> 6513

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<221> misc_feature
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<210> 6514
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<221> misc_feature
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<223> Accession number cg29242150

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<210> 6519
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<221> misc_feature
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<223> Accession number cg29242150

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<210> 6520
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<223> Accession number cg29242150

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<210> 6521
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<221> misc_feature
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<223> Accession number cg39667665

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<210> 6522
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<210> 6524
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<223> 2 of 2 allelic variants (6525 is other entry)

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<223> Accession number cg39667665

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<212> DNA

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<222> (0)...(0)

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51

<210> 6528

<211> 51

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<213> Homo sapiens

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<223> 2 of 2 allelic variants (6527 is other entry)

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<223> Accession number cg43261509

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51

<210> 6529

<211> 51

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<221> misc_feature

<222> (0)...(0)

<223> Accession number cg38629253

<400> 6529
ccaccacact cggcctccca aagtgttggg attacaggtg tgagccaccg g 51

<210> 6530
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 2 of 2 allelic variants (6529 is other entry)

<221> misc_feature
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<223> Accession number cg38629253

<400> 6530
ccaccacact cggcctccca aagtgttggg attacaggtg tgagccaccg g 51

<210> 6531
<211> 51
<212> DNA
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<223> 1 of 2 allelic variants (6532 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg39710199

<400> 6531
tgatcccgat gggaatgacg agagctagta gcagctgctc accgttgcg a 51

<210> 6532
<211> 51
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<213> Homo sapiens

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<223> 2 of 2 allelic variants (6531 is other entry)

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<223> Accession number cg39710199

<400> 6532
tgatcccgat gggaatgacg agagccagta gcagctgctc accgttgcg a 51

<210> 6533
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<213> Homo sapiens

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<222> (26)...(0)

<223> 1 of 2 allelic variants (6534 is other entry)

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<222> (0)...(0)

<223> Accession number cg29219974

<400> 6533

acgacggcgg ttaccgcccc gaccgaggct ggaacaagtg ttcccgctc g

51

<210> 6534

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 2 of 2 allelic variants (6533 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg29219974

<400> 6534

acgacggcgg ttaccgcccc gaccgtggct ggaacaagtg ttcccgctc g

51

<210> 6535

<211> 51

<212> DNA

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<223> 1 of 2 allelic variants (6536 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg29219974

<400> 6535

gttaccgccc cgaccgggc tgaacaagt gttcccgct cgtcggtag g

51

<210> 6536

<211> 51

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<223> 2 of 2 allelic variants (6535 is other entry)

<221> misc_feature
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<223> Accession number cg29219974

<400> 6536
gttacgcgcc cgaccgcggc tggaataagt gttcccgctg cgtcggtgac g 51

<210> 6537
<211> 51
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<223> 1 of 2 allelic variants (6538 is other entry)

<221> misc_feature
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<223> Accession number cg38821538

<400> 6537
aaaaattagc caggcgtggt ggtacacgcc tgtaatccca gctactcagg a 51

<210> 6538
<211> 51
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<213> Homo sapiens

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<223> 2 of 2 allelic variants (6537 is other entry)

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<223> Accession number cg38821538

<400> 6538
aaaaattagc caggcgtggt ggtacgcgcc tgtaatccca gctactcagg a 51

<210> 6539
<211> 51
<212> DNA
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<223> 1 of 2 allelic variants (6540 is other entry)

<221> misc_feature
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<400> 6539

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51

<210> 6540

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 2 of 2 allelic variants (6539 is other entry)

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<223> Accession number cg38821538

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51

<210> 6541

<211> 51

<212> DNA

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<223> 1 of 2 allelic variants (6542 is other entry)

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<222> (0)...(0)

<223> Accession number cg43297632

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51

<210> 6542

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 2 of 2 allelic variants (6541 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43297632

<400> 6542

catggtgaaa ccctgtctct actaagaaaa tacagaaaat tagctgggcg t

51

<210> 6543

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 1 of 2 allelic variants (6544 is other entry)

<221> misc_feature
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<223> Accession number cg43297632

<400> 6543
aaattagctg ggcgtggtgg cgggcgcctg tagccccagc tacttgggag g 51

<210> 6544
<211> 51
<212> DNA
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<223> 2 of 2 allelic variants (6543 is other entry)

<221> misc_feature
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<223> Accession number cg43297632

<400> 6544
aaattagctg ggcgtggtgg cgggcacctg tagccccagc tacttgggag g 51

<210> 6545
<211> 51
<212> DNA
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<220>
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<223> 1 of 2 allelic variants (6546 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg33204852

<400> 6545
cttcgagggc gtcacggtga tcaagctggg gaagaatccc aggtaaccgt c 51

<210> 6546
<211> 51
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<223> 2 of 2 allelic variants (6545 is other entry)

<221> misc_feature
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<223> Accession number cg33204852

<400> 6546
cttcgagggc gtcacggtga tcaagttggg gaagaatccc aggtaaccgt c 51

<210> 6547
<211> 51
<212> DNA
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<223> 1 of 2 allelic variants (6548 is other entry)

<221> misc_feature
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<223> Accession number cg43999183

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gtgaggcagg agaatcgctt gaacctagga ggcagaggct gcagtgggcc g 51

<210> 6548
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<223> 2 of 2 allelic variants (6547 is other entry)

<221> misc_feature
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<223> Accession number cg43999183

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gtgaggcagg agaatcgctt gaacccagga ggcagaggct gcagtgggcc g 51

<210> 6549
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<223> 1 of 2 allelic variants (6550 is other entry)

<221> misc_feature
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<223> Accession number cg43982145

<400> 6549
tcaggagac gcggtaaaag tagttcctcc agaacacttc ttccttcaca a 51

<210> 6550
<211> 51
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<223> 2 of 2 allelic variants (6549 is other entry)

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<223> Accession number cg43982145

<400> 6550
tcagggagac gcggtaaaag tagtttctcc agaacacttc ttccttcaca a 51

<210> 6551
<211> 51
<212> DNA
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<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (6552 is other entry)

<221> misc_feature
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<223> Accession number cg21632104

<400> 6551
ggtcgatggc gtcgacagcc cgttctggac ggggccagaa cagccgattt c 51

<210> 6552
<211> 51
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<222> (26)...(0)
<223> 2 of 2 allelic variants (6551 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg21632104

<400> 6552
ggtcgatggc gtcgacagcc cgttccggac ggggccagaa cagccgattt c 51

<210> 6553
<211> 51
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<213> Homo sapiens

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<222> (26)...(0)
<223> 1 of 2 allelic variants (6554 is other entry)

<221> misc_feature
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<223> Accession number cg21632104

<400> 6553
tggcgtcgac agcccgttct ggacgggtcc agaacagccg atttccgcga c 51

<210> 6554
<211> 51
<212> DNA
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<223> 2 of 2 allelic variants (6553 is other entry)

<221> misc_feature
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<400> 6554
tggcgtcgac agcccgttct ggacgagtcc agaacagccg atttccgcga c 51

<210> 6555
<211> 51
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<223> 1 of 2 allelic variants (6556 is other entry)

<221> misc_feature
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<223> Accession number cg21632104

<400> 6555
gcggggccgag cacatacggg ttatccgggt gcaacacagt cgattccacc g 51

<210> 6556
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (6555 is other entry)

<221> misc_feature

<222> (0)...(0)
<223> Accession number cg21632104

<400> 6556
gcggggccgag cacatacggg ttatctgggt gcaacacagt cgattccacc g 51

<210> 6557
<211> 51
<212> DNA
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<222> (26)...(0)
<223> 1 of 2 allelic variants (6558 is other entry)

<221> misc_feature
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<223> Accession number cg21632104

<400> 6557
agcacatacg gggttatccgg gtgcaacaca gtcgattcca ccgaggacga g 51

<210> 6558
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 2 of 2 allelic variants (6557 is other entry)

<221> misc_feature
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<223> Accession number cg21632104

<400> 6558
agcacatacg gggttatccgg gtgcagcaca gtcgattcca ccgaggacga g 51

<210> 6559
<211> 51
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<222> (26)...(0)
<223> 1 of 2 allelic variants (6560 is other entry)

<221> misc_feature
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<223> Accession number cg21632104

<400> 6559
acgggttatc cgggtgcaac acagtcgatt ccaccgagga cgagaacaat a 51

<210> 6560
<211> 51
<212> DNA
<213> Homo sapiens

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<222> (26)...(0)
<223> 2 of 2 allelic variants (6559 is other entry)

<221> misc_feature
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<223> Accession number cg21632104

<400> 6560
acgggttatc cgggtgcaac acagttgatt ccaccgagga cgagaacaat a 51

<210> 6561
<211> 51
<212> DNA
<213> Homo sapiens

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<222> (26)...(0)
<223> 1 of 2 allelic variants (6562 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg21632104

<400> 6561
ccgggttgta ttcgtcgacc agccattgat cccctgatg caggtacacc g 51

<210> 6562
<211> 51
<212> DNA
<213> Homo sapiens

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<222> (26)...(0)
<223> 2 of 2 allelic variants (6561 is other entry)

<221> misc_feature
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<223> Accession number cg21632104

<400> 6562
ccgggttgta ttcgtcgacc agccactgat cccctgatg caggtacacc g 51

<210> 6563
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<212> DNA
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<222> (26)...(0)
<223> 1 of 2 allelic variants (6564 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg21659365

<400> 6563
ggattggcat cttcattggc gctctcacct tcacggggtc gctggtggcc t 51

<210> 6564
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 2 of 2 allelic variants (6563 is other entry)

<221> misc_feature
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<223> Accession number cg21659365

<400> 6564
ggattggcat cttcattggc gctcttacct tcacggggtc gctggtggcc t 51

<210> 6565
<211> 51
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<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (6566 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg21659365

<400> 6565
tcattggcgc tctcaccttc acggggtcgc tgggtggcctg gggcaagctc t 51

<210> 6566
<211> 51
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<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (6565 is other entry)

<221> misc_feature
<222> (0)...(0)

<223> Accession number cg21659365

<400> 6566

tcattggcgc tctcaccttc acgggatcgc tggtaggctg gggcaagctc t

51

<210> 6567

<211> 51

<212> DNA

<213> Homo sapiens

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<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (6568 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg44126917

<400> 6567

cctgtgggac ctttccatca gcagccagcg tcgctgcagc aatggccaga t

51

<210> 6568

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (6567 is other entry)

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<222> (0)...(0)

<223> Accession number cg44126917

<400> 6568

cctgtgggac ctttccatca gcagcgagcg tcgctgcagc aatggccaga t

51

<210> 6569

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (6570 is other entry)

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<222> (0)...(0)

<223> Accession number cg44126917

<400> 6569

caaaatgcgt cccagccttt ggtaggccag ctggcgtgag attgatcgta a

51

<210> 6570

<211> 51
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<222> (26)...(0)
<223> 2 of 2 allelic variants (6569 is other entry)

<221> misc_feature
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<223> Accession number cg44126917

<400> 6570
caaatgcgt cccagccttt ggtagaccag ctggcgtgag attgatcgtc a 51

<210> 6571
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (6572 is other entry)

<221> misc_feature
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<223> Accession number cg42286566

<400> 6571
caatttggtc tcgaactcct gagctcaagt gatccgcccgc ccttggcctc c 51

<210> 6572
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 2 of 2 allelic variants (6571 is other entry)

<221> misc_feature
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<223> Accession number cg42286566

<400> 6572
caatttggtc tcgaactcct gagcttaagt gatccgcccgc ccttggcctc c 51

<210> 6573
<211> 51
<212> DNA
<213> Homo sapiens

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<222> (26)...(0)
<223> 1 of 2 allelic variants (6574 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43926000

<400> 6573
aggtcaggag ttcaagacca gcctagccaa aatggtgaaa ccccgctctct a 51

<210> 6574
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (6573 is other entry)

<221> misc_feature
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<223> Accession number cg43926000

<400> 6574
aggtcaggag ttcaagacca gcctaacc aa aatggtgaaa ccccgctctct a 51

<210> 6575
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (6576 is other entry)

<221> misc_feature
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<223> Accession number cg43967015

<400> 6575
gggaggctga ggcaggcgga tcacctgaga ttaggagttc gagactagcc t 51

<210> 6576
<211> 51
<212> DNA
<213> Homo sapiens

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<222> (26)...(0)
<223> 2 of 2 allelic variants (6575 is other entry)

<221> misc_feature
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<223> Accession number cg43967015

<400> 6576
gggaggctga ggcaggcgga tcacccgaga ttaggagttc gagactagcc t 51

<210> 6577
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (6578 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43967015

<400> 6577
ccatctctac taaaaatata aaaattagcc aggcgtggtg gcgcatgcct g 51

<210> 6578
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 2 of 2 allelic variants (6577 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43967015

<400> 6578
ccatctctac taaaaataca aaaatcagcc aggcgtggtg gcgcatgcct g 51

<210> 6579
<211> 51
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (26)...(0)
<223> 1 of 2 allelic variants (6580 is other entry)

<221> misc_feature
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<223> Accession number cg43967015

<400> 6579
ggtggcgcat gcctgtaatc tcagctactc gggaggctga ggcaggagaa t 51

<210> 6580
<211> 51

2008

<223> 1 of 2 allelic variants (6584 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg20729392

<400> 6583

cgatggcagt caactcagcg atgcccgatg tcatccagtt cagcatgtac a

51

<210> 6584

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (6583 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg20729392

<400> 6584

cgatggcagt caactcagcg atgcctgatg tcatccagtt cagcatgtac a

51

<210> 6585

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 1 of 2 allelic variants (6586 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg20729392

<400> 6585

cccgggcata ggacacaaaa gaaccactag aggggcggtg aatgaccaac t

51

<210> 6586

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (6585 is other entry)

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<223> Accession number cg20729392

<400> 6586
cccgggcata ggacacaaaa gaaccgctag aggggcggtg aatgaccaac t 51

<210> 6587
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<222> (26)...(0)
<223> 1 of 2 allelic variants (6588 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43273813

<400> 6587
aacatggtga aaccccgctct ctactaaaaa tataaaaaat tagccaggca t 51

<210> 6588
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (6587 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43273813

<400> 6588
aacatggtga aaccccgctct ctactgaaaa tataaaaaat tagccaggca t 51

<210> 6589
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (6590 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg42487874

<400> 6589
aaaaaaaatt agccagggtgt ggtggtgggc gcctgtggtc cgggctgctc g 51

<210> 6590
<211> 51
<212> DNA

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<223> 1 of 2 allelic variants (6592 is other entry)

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<223> Accession number cg44019253

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<210> 6592

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<223> Accession number cg44019253

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<210> 6595
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<210> 6596
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<210> 6597

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<210> 6598

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<223> Accession number cg29344427

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51

<210> 6599

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51

<210> 6600

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<210> 6601
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<210> 6602
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<210> 6603
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<210> 6604
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<210> 6605
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<210> 6606
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51

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51

<210> 6610

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<210> 6612
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<210> 6614
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<210> 6616
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<210> 6618
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<210> 6625

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51

<210> 6626

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<210> 6632
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<210> 6633
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<210> 6634
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<210> 6636
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<210> 6638
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<223> Accession number cg21632104

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<210> 6645
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<210> 6646
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<210> 6647
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<223> Accession number cg37854509

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51

<210> 6648

<211> 51

<212> DNA

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<223> 2 of 2 allelic variants (6647 is other entry)

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<223> Accession number cg37854509

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51

<210> 6649

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<212> DNA

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<223> 1 of 2 allelic variants (6650 is other entry)

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51

<210> 6650

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 2 of 2 allelic variants (6649 is other entry)

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<210> 6651
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tcggctacga gtactctccg gagatcttta gccagaccg cacggacttc g 51

<210> 6652
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<210> 6653
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taatctgccg gctaccgtcg agatggggcac tccgaacacc tacgccgacc a 51

<210> 6654

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<223> 2 of 2 allelic variants (6653 is other entry)

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<223> Accession number cg32128336

<400> 6654

taatctgccg gctaccgtcg agatgagcac tccgaacacc tacgccgacc a 51

<210> 6655

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 1 of 2 allelic variants (6656 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg38912931

<400> 6655

taagttcact ggcctctggg agtctggccc tgaggaccaa ctaacaacac c 51

<210> 6656

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 2 of 2 allelic variants (6655 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg38912931

<400> 6656

taagttcact ggcctctggg agtcttgccc tgaggaccaa ctaacaacac c 51

<210> 6657

<211> 51

<212> DNA

<213> Homo sapiens

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<221> misc_feature
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<210> 6658
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<223> 2 of 2 allelic variants (6657 is other entry)

<221> misc_feature
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<400> 6658
ggacccttag caaaactact gtctgcagtg aaagaaatct ctggctctaa g 51

<210> 6659
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<212> DNA
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<220>
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<223> 1 of 2 allelic variants (6660 is other entry)

<221> misc_feature
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<223> Accession number cg42512386

<400> 6659
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<210> 6660
<211> 51
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<213> Homo sapiens

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<210> 6661
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<223> Accession number cg42719442

<400> 6661
agcaaataca cccaggagga gtagaggga ggaaataatc aaactcagag c 51

<210> 6662
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<223> 2 of 2 allelic variants (6661 is other entry)

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<400> 6662
agcaaataca cccaggagga gtagaaggca ggaaataatc aaactcagag c 51

<210> 6663
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<223> 1 of 2 allelic variants (6664 is other entry)

<221> misc_feature
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<223> Accession number cg43974178

<400> 6663
tgtgattgtg gaggaacaga cagaggagac ccaagtgact gaagaagtga c 51

<210> 6664
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<223> 2 of 2 allelic variants (6663 is other entry)

<221> misc_feature
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<223> Accession number cg43974178

<400> 6664
tgtgattgtg gaggaacaga cagagaagac ccaagtgact gaagaagtga c 51

<210> 6665
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<223> 1 of 2 allelic variants (6666 is other entry)

<221> misc_feature
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<223> Accession number cg38335527

<400> 6665
acggcgtgca aggacaagca gctgcagatc tttgacccca gaacaaagcc g 51

<210> 6666
<211> 51
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<223> 2 of 2 allelic variants (6665 is other entry)

<221> misc_feature
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<400> 6666
acggcgtgca aggacaagca gctgcggatc tttgacccca gaacaaagcc g 51

<210> 6667
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<223> 1 of 2 allelic variants (6668 is other entry)

<221> misc_feature
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<223> Accession number cg39517733

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cacttggggt agcagccttc caggctacgg agatcttgta ggcactgtcc a 51

<210> 6668
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 2 of 2 allelic variants (6667 is other entry)

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<223> Accession number cg39517733

<400> 6668
cacttggggt agcagccttc caggccacgg agatcttgta ggcactgtcc a 51

<210> 6669
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<223> 1 of 2 allelic variants (6670 is other entry)

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<223> Accession number cg44127439

<400> 6669
gcctccttga tggaagccgc cttcttgctg tcgacaaacc ccaaacagac a 51

<210> 6670
<211> 51
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<223> 2 of 2 allelic variants (6669 is other entry)

<221> misc_feature

<222> (0)...(0)
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<400> 6670
gcctccttga tggaagccgc cttctcgtcg tcgacaaacc ccaaacagac a 51

<210> 6671
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<223> 1 of 2 allelic variants (6672 is other entry)

<221> misc_feature
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<400> 6671
tcgtaggagt gggtagtgca gatctcgccg taatgagagc gagcgggtatt g 51

<210> 6672
<211> 51
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<223> 2 of 2 allelic variants (6671 is other entry)

<221> misc_feature
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<400> 6672
tcgtaggagt gggtagtgca gatcttgccg taatgagagc gagcgggtatt g 51

<210> 6673
<211> 51
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<223> 1 of 2 allelic variants (6674 is other entry)

<221> misc_feature
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<223> Accession number cg43103077

<400> 6673
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<210> 6674
<211> 51
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<223> 2 of 2 allelic variants (6673 is other entry)

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<223> Accession number cg43103077

<400> 6674
gaatgagctg tgtctaccag ttcttttagc tcttcagggtg ttccataacg g

51

<210> 6675
<211> 51
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<223> 1 of 2 allelic variants (6676 is other entry)

<221> misc_feature
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<223> Accession number cg43103077

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cctttgattc ttggtagtagc attgcatgta aaatgtttat aagaagctac t

51

<210> 6676
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<221> misc_feature
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<400> 6676
cctttgattc ttggtagtagc attgcttgta aaatgtttat aagaagctac t

51

<210> 6677
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<223> 1 of 2 allelic variants (6678 is other entry)

<221> misc_feature
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cttagactcc gtggcttctt tggctctggaa tgcttaaact catatgagtg t 51

<210> 6678
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<223> 2 of 2 allelic variants (6677 is other entry)

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<400> 6678
cttagactcc gtggcttctt tggctcggaa tgcttaaact catatgagtg t 51

<210> 6679
<211> 51
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<213> Homo sapiens

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<223> 1 of 2 allelic variants (6680 is other entry)

<221> misc_feature
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<223> Accession number cg39433010

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<210> 6680
<211> 51
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<213> Homo sapiens

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<223> 2 of 2 allelic variants (6679 is other entry)

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<223> Accession number cg39433010

<400> 6680

tctaattgtac cttcctgcat agttatctag ttgggtaatg ggaggttgga a

51

<210> 6681

<211> 51

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<213> Homo sapiens

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<222> (26)...(0)

<223> 1 of 2 allelic variants (6682 is other entry)

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<222> (0)...(0)

<223> Accession number cg38434693

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51

<210> 6682

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 2 of 2 allelic variants (6681 is other entry)

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<222> (0)...(0)

<223> Accession number cg38434693

<400> 6682

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51

<210> 6683

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 1 of 2 allelic variants (6684 is other entry)

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51

<210> 6684

<211> 51
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<223> 2 of 2 allelic variants (6683 is other entry)

<221> misc_feature
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<210> 6685
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<223> 1 of 2 allelic variants (6686 is other entry)

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<210> 6686
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<220>
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<223> 2 of 2 allelic variants (6685 is other entry)

<221> misc_feature
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gcctctgggt tgtacttata atccattctc agatctgtgg tgctggcact g 51

<210> 6687
<211> 51
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<223> 1 of 2 allelic variants (6688 is other entry)

<221> misc_feature
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<223> Accession number cg39513248

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gcagcttgca agatcgggtgt tagacgggga ttcagtcaga gccaggtgga a 51

<210> 6688
<211> 51
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<223> 2 of 2 allelic variants (6687 is other entry)

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<400> 6688
gcagcttgca agatcgggtgt tagactggga ttcagtcaga gccaggtgga a 51

<210> 6689
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<221> misc_feature
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<400> 6689
agaggataat aaagaacatt catttgaggt ttcattgttt gcggaacttt t 51

<210> 6690
<211> 51
<212> DNA
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<223> 2 of 2 allelic variants (6689 is other entry)

<221> misc_feature
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<400> 6690
agaggataat aaagaacatt catttttaggt ttcattgttt gcggaacttt t 51

<210> 6691
<211> 51
<212> DNA
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<223> 1 of 2 allelic variants (6692 is other entry)

<221> misc_feature
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<223> Accession number cg43287601

<400> 6691
ttgaactcct gacctcaagt gatccgcccg cctcggcctc ccaaagtgt g 51

<210> 6692
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (6691 is other entry)

<221> misc_feature
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<223> Accession number cg43287601

<400> 6692
ttgaactcct gacctcaagt gatccaccg cctcggcctc ccaaagtgt g 51

<210> 6693
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<223> 1 of 2 allelic variants (6694 is other entry)

<221> misc_feature
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<223> Accession number cg43967859

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<210> 6694
<211> 51

<212> DNA
<213> Homo sapiens

<220>
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<223> 2 of 2 allelic variants (6693 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg43967859

<400> 6694
gtgtgtatca gctggccctt aaatctctgg agaaaaccat cttcattaaa a 51

<210> 6695
<211> 51
<212> DNA
<213> Homo sapiens

<220>
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<222> (26)...(0)
<223> 1 of 2 allelic variants (6696 is other entry)

<221> misc_feature
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<223> Accession number cg39527289

<400> 6695
cccgtcacc gaggacaaaa ctgttcgcga gaacgtcgaa gaggccgtcg g 51

<210> 6696
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 2 of 2 allelic variants (6695 is other entry)

<221> misc_feature
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<400> 6696
cccgtcacc gaggacaaaa ctgtttgcga gaacgtcgaa gaggccgtcg g 51

<210> 6697
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<212> DNA
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<223> 1 of 2 allelic variants (6698 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43923983

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51

<210> 6698

<211> 51

<212> DNA

<213> Homo sapiens

<220>

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<222> (26)...(0)

<223> 2 of 2 allelic variants (6697 is other entry)

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<223> Accession number cg43923983

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51

<210> 6699

<211> 51

<212> DNA

<213> Homo sapiens

<220>

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<222> (26)...(0)

<223> 1 of 2 allelic variants (6700 is other entry)

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<222> (0)...(0)

<223> Accession number cg42920024

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caatgagccg agatcatgcc actgcgctcc agcctgggca acagggtgag a

51

<210> 6700

<211> 51

<212> DNA

<213> Homo sapiens

<220>

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<223> 2 of 2 allelic variants (6699 is other entry)

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<222> (0)...(0)

<223> Accession number cg42920024

<400> 6700
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<210> 6701
<211> 51
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<223> 1 of 2 allelic variants (6702 is other entry)

<221> misc_feature
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<223> Accession number cg43302383

<400> 6701
ctgaaccac aggacttcat tggctgcctg aacgtgaagg cgacttttta t 51

<210> 6702
<211> 51
<212> DNA
<213> Homo sapiens

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<210> 6703
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<212> DNA
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<222> (26)...(0)
<223> 1 of 2 allelic variants (6704 is other entry)

<221> misc_feature
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<223> Accession number cg42734969

<400> 6703
aagtgcacaa ctctctgtgc tgatgacat gatgagaaat atggagtccc t 51

<210> 6704
<211> 51
<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (26)...(0)

<223> 2 of 2 allelic variants (6703 is other entry)

<221> misc_feature

<222> (0)...(0)

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<400> 6704

aagtgcacaa ctctctgtgc tgatggccat gatgagaaat atggagtccc t

51

<210> 6705

<211> 51

<212> DNA

<213> Homo sapiens

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<222> (26)...(0)

<223> 1 of 2 allelic variants (6706 is other entry)

<221> misc_feature

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51

<210> 6706

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<223> Accession number cg43989360

<400> 6706

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51

<210> 6707

<211> 51

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<223> 1 of 2 allelic variants (6708 is other entry)

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<210> 6708
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<210> 6709
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<210> 6710
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51

<210> 6711

<211> 51

<212> DNA

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<223> 1 of 2 allelic variants (6712 is other entry)

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<400> 6711

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<210> 6712

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<212> DNA

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<223> 1 of 2 allelic variants (6714 is other entry)

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<223> Accession number cg43280136

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51

<210> 6714

<211> 51

<212> DNA

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<210> 6715
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<210> 6716
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<210> 6717
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<210> 6718
<211> 51
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<210> 6719
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<223> 1 of 2 allelic variants (6720 is other entry)

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<210> 6720
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<221> misc_feature
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<210> 6721
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<223> Accession number cg44127437

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<210> 6722
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<223> 2 of 2 allelic variants (6721 is other entry)

<221> misc_feature
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<210> 6723
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<223> 1 of 2 allelic variants (6724 is other entry)

<221> misc_feature
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<223> Accession number cg42868441

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tcccagctac ttgggaggct gaggcaggag aattgcttga acccaggagg c 51

<210> 6724
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<210> 6725
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<223> 1 of 2 allelic variants (6726 is other entry)

<221> misc_feature
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<223> Accession number cg29256713

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cgattcggtg acgctgagtc gtttctgaac atcatcgatt ccattcgctc t 51

<210> 6726
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<223> 2 of 2 allelic variants (6725 is other entry)

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cgattcggtg acgctgagtc gtttccgaac atcatcgatt ccattcgctc t 51

<210> 6727
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<223> 1 of 2 allelic variants (6728 is other entry)

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<222> (0)...(0)
<223> Accession number cg43921362

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atggacgtta actgacctaa atacttaacc tccttctcaa tatagtcgct c 51

<210> 6728
<211> 51
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<223> 2 of 2 allelic variants (6727 is other entry)

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atggacgtta actgacctaa atactcaacc tccttctcaa tatagtcgct c 51

<210> 6729
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<223> 1 of 2 allelic variants (6730 is other entry)

<221> misc_feature
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<223> Accession number cg38213630

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<210> 6730
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<223> 2 of 2 allelic variants (6729 is other entry)

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<223> Accession number cg38213630

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<210> 6731
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<223> Accession number cg41616497

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<210> 6732
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<210> 6733
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<221> misc_feature
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<223> Accession number cg43320455

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<210> 6734
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<221> misc_feature
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cttgttcttt acgatgtagg catcctctgg ttccagcagg aagtgaggca g 51

<210> 6736
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<223> 2 of 2 allelic variants (6735 is other entry)

<221> misc_feature
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<210> 6737
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<223> 1 of 2 allelic variants (6738 is other entry)

<221> misc_feature
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<223> Accession number cg43929320

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gaagatctct ctgcaagagt agatgcagtt aaggaagaaa atctgaagct a

51

<210> 6738

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 2 of 2 allelic variants (6737 is other entry)

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<223> Accession number cg43929320

<400> 6738

gaagatctct ctgcaagagt agatgaagtt aaggaagaaa atctgaagct a

51

<210> 6739

<211> 51

<212> DNA

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<223> 1 of 2 allelic variants (6740 is other entry)

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<223> Accession number cg35929441

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51

<210> 6740

<211> 51

<212> DNA

<213> Homo sapiens

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51

<210> 6741

<211> 51
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<221> misc_feature
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<223> Accession number cg35929441

<400> 6741
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<210> 6742
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<212> DNA
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<223> 2 of 2 allelic variants (6741 is other entry)

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<223> Accession number cg35929441

<400> 6742
tttccttaca ccatattaca aaaatcaact caagatggat taaagactta a 51

<210> 6743
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<221> misc_feature
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<400> 6743
gggcaaagat ttcattgacaa aaacacacaaa agcaattgca acaaaagcaa a 51

<210> 6744
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<222> (26)...(0)
<223> 2 of 2 allelic variants (6743 is other entry)

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gggcaaagat ttcattgacaa aaacatcaaa agcaattgca acaaaagcaa a 51

<210> 6745
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<223> 1 of 2 allelic variants (6746 is other entry)

<221> misc_feature
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<223> Accession number cg42667624

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<210> 6746
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<221> misc_feature
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<223> Accession number cg42667624

<400> 6746
ctgggtctgta tatggttggt aattcagtggt ctccatgggg taacaagggc c 51

<210> 6747
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<221> misc_feature
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<223> Accession number cg39667665

<400> 6747
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<210> 6748
<211> 51
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<223> 2 of 2 allelic variants (6747 is other entry)

<221> misc_feature
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<400> 6748
atcccagcac tttgggaggc cgagggtgggc ggatcacctg aggtcaggag t 51

<210> 6749
<211> 51
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<223> 1 of 2 allelic variants (6750 is other entry)

<221> misc_feature
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<223> Accession number cg42481380

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ggagaatggt gtgaaccagc gaggcggagc ttgcagtgcg ccaatatcgc g 51

<210> 6750
<211> 51
<212> DNA
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<223> 2 of 2 allelic variants (6749 is other entry)

<221> misc_feature
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<223> Accession number cg42481380

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ggagaatggt gtgaaccagc gaggcagagc ttgcagtgcg ccaatatcgc g 51

<210> 6751
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<212> DNA
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<223> 1 of 2 allelic variants (6752 is other entry)

<221> misc_feature
<222> (0)...(0)
<223> Accession number cg42208556

<400> 6751
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51

<210> 6752
<211> 51
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<223> 2 of 2 allelic variants (6751 is other entry)

<221> misc_feature
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<400> 6752
agaccagcct ggccaatgta gcgaagcccc atctctacta caaatataaa a

51

<210> 6753
<211> 51
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<222> (26)...(0)
<223> 1 of 2 allelic variants (6754 is other entry)

<221> misc_feature
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<223> Accession number cg43261509

<400> 6753
ctttgggagg ctgaggcggg cggatcactt aaggtcagga gttcaagacc a

51

<210> 6754
<211> 51
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<213> Homo sapiens

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<223> 2 of 2 allelic variants (6753 is other entry)

<221> misc_feature

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<223> Accession number cg43261509

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ctttgggagg ctgaggcggg cggattactt aaggtcagga gttcaagacc a

51

<210> 6755

<211> 51

<212> DNA

<213> Homo sapiens

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<221> misc_feature

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<223> 1 of 2 allelic variants (6756 is other entry)

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<222> (0)...(0)

<223> Accession number cg25239778

<400> 6755

aaccgggaa gtggaggctg cagtgagcca ggatcatgcc actgcactcc a

51

<210> 6756

<211> 51

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<223> 2 of 2 allelic variants (6755 is other entry)

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<223> Accession number cg25239778

<400> 6756

aaccgggaa gtggaggctg cagtgaggcca ggatcatgcc actgcactcc a

51

<210> 6757

<211> 51

<212> DNA

<213> Homo sapiens

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<221> misc_feature

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<223> 1 of 2 allelic variants (6758 is other entry)

<221> misc_feature

<222> (0)...(0)

<223> Accession number cg38629253

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<210> 6758
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<212> DNA
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<223> 2 of 2 allelic variants (6757 is other entry)

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<210> 6759
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<221> misc_feature
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<210> 6760
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<210> 6761
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<213> Homo sapiens

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<223> 1 of 2 allelic variants (6762 is other entry)

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51

<210> 6762

<211> 51

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<223> 2 of 2 allelic variants (6761 is other entry)

<221> misc_feature

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<223> Accession number cg39710199

<400> 6762

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51

<210> 6763

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<212> DNA

<213> Homo sapiens

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<223> Accession number cg43280209

<400> 6763

gactacagat gccacaacc acgcctagct aatttttgta ttttttagtag a

51

<210> 6764

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 2 of 2 allelic variants (6763 is other entry)

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<210> 6765
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<223> 1 of 2 allelic variants (6766 is other entry)

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cctgatgggc tagcccatc tgacttgatg tgtacagggg ataaggacgt g 51

<210> 6766
<211> 51
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<223> 2 of 2 allelic variants (6765 is other entry)

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cctgatgggc tagcccatc tgactcgatg tgtacagggg ataaggacgt g 51

<210> 6767
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<223> Accession number cg38821538

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gttcaagatc agcctggcca acatgatgaa accctatctc tactaaaatt a 51

<210> 6768

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<223> 2 of 2 allelic variants (6767 is other entry)

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<223> Accession number cg38821538

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gttcaagatc agcctggcca acatggtgaa accctatctc tactaaaatt a 51

<210> 6769

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 1 of 2 allelic variants (6770 is other entry)

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<223> Accession number cg38821538

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<210> 6770

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 2 of 2 allelic variants (6769 is other entry)

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<223> Accession number cg38821538

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tacacgctg taatcccagc tactcgggag gctgaggcag gagaattggt t 51

<210> 6771

<211> 51

<212> DNA

<213> Homo sapiens

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<223> 1 of 2 allelic variants (6772 is other entry)

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<400> 6771
gccatgcctt gtcttttgggt ctcataaata gtcactgggg ccgggcgcag t 51

<210> 6772
<211> 51
<212> DNA
<213> Homo sapiens

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<223> Accession number cg43297632

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gccatgcctt gtcttttgggt ctcataaata gtcactgggg ccgggcgcag t 51

<210> 6773
<211> 51
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<223> Accession number cg43297632

<400> 6773
ataaatagtc actggggccg ggcgcagtga ctcacgcctg taatcccagc a 51

<210> 6774
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 2 of 2 allelic variants (6773 is other entry)

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<223> Accession number cg43297632

<400> 6774
ataaatagtc actggggccg ggcgcggtga ctcacgcctg taatcccagc a 51

<210> 6775
<211> 51
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<210> 6776
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gcggggcgct gtagccccag ctactcggga ggctgaggcg ggagaatggc a 51

<210> 6777
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<223> 1 of 2 allelic variants (6778 is other entry)

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<400> 6777
gccccagcta cttgggaggc tgaggcggga gaatggcaat ggcgtgaacc c 51

<210> 6778
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 2 of 2 allelic variants (6777 is other entry)

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<400> 6778
gccccagcta cttgggaggc tgaggtggga gaatggcaat ggcgtaacc c 51

<210> 6779
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<223> 1 of 2 allelic variants (6780 is other entry)

<221> misc_feature
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<223> Accession number cg42657675

<400> 6779
ggctcatgcc tgtaatccca gcattttggg aggccgagat gggcggatca c 51

<210> 6780
<211> 51
<212> DNA
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<223> 2 of 2 allelic variants (6779 is other entry)

<221> misc_feature
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<223> Accession number cg42657675

<400> 6780
ggctcatgcc tgtaatccca gcattctggg aggccgagat gggcggatca c 51

<210> 6781
<211> 51
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<223> 1 of 2 allelic variants (6782 is other entry)

<221> misc_feature
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<223> Accession number cg23333150

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cgcccgcag catgcccggc caattctttg tatttttagt agagacgggg t 51

<210> 6782
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<212> DNA
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<223> 2 of 2 allelic variants (6781 is other entry)

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cgcccgcag catgcccggc caattttttg tatttttagt agagacgggg t 51

<210> 6783
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<221> misc_feature
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<223> Accession number cg42475469

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tcaccatggt gccagactg gtctcgaact cctgacctca agtgatccac c 51

<210> 6784
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<223> 2 of 2 allelic variants (6783 is other entry)

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<222> (0)...(0)
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tcaccatggt ggccagactg gtctcaaact cctgacctca agtgatccac c 51

<210> 6785
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<223> 1 of 2 allelic variants (6786 is other entry)

<221> misc_feature
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<223> Accession number cg44126917

<400> 6785
tttccatcag cagccagcgt cgctgcagca atggccagat caaaatgcgt c 51

<210> 6786
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<223> 2 of 2 allelic variants (6785 is other entry)

<221> misc_feature
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<210> 6787
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<223> 1 of 2 allelic variants (6788 is other entry)

<221> misc_feature
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<223> Accession number cg44126917

<400> 6787
agttctccaa tcagaaccgt cgatcccaag agagcctgtg ggacctttcc a 51

<210> 6788
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<212> DNA
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<223> 2 of 2 allelic variants (6787 is other entry)

<221> misc_feature
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<223> Accession number cg44126917

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agttctccaa tcagaaccgt cgatctcaag agagcctgtg ggaccttcc a 51

<210> 6789
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<223> 1 of 2 allelic variants (6790 is other entry)

<221> misc_feature
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<400> 6789
aatttggtct cgaactcctg agctcaagtg atccgccgc cttggcctcc c 51

<210> 6790
<211> 51
<212> DNA
<213> Homo sapiens

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<223> 2 of 2 allelic variants (6789 is other entry)

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<223> Accession number cg42286566

<400> 6790
aatttggtct cgaactcctg agctcgagt atccgccgc cttggcctcc c 51

<210> 6791
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<223> 1 of 2 allelic variants (6792 is other entry)

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<223> Accession number cg43926000

<400> 6791
taatcccagc actttgggag gccaaaggtgg gcggatcact tgaggtcagg a 51

<210> 6792
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<210> 6793
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<223> 1 of 2 allelic variants (6794 is other entry)

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gcactttggg aggccaaaggt gggcggatca cttgaggtca ggagttcaag a 51

<210> 6794
<211> 51
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<223> 2 of 2 allelic variants (6793 is other entry)

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<223> Accession number cg43926000

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51

<210> 6795

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<212> DNA

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<223> 1 of 2 allelic variants (6796 is other entry)

<221> misc_feature

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<223> Accession number cg43926000

<400> 6795

tcacttgagg tcaggagttc aagaccagcc tagccaaaat ggtgaaaccc c

51

<210> 6796

<211> 51

<212> DNA

<213> Homo sapiens

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<221> misc_feature

<222> (0)...(0)

<223> Accession number cg43926000

<400> 6796

tcacttgagg tcaggagttc aagactagcc tagccaaaat ggtgaaaccc c

51

<210> 6797

<211> 51

<212> DNA

<213> Homo sapiens

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<223> Accession number cg43967015

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51

<210> 6798

2073

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<221> misc_feature
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<223> Accession number cg43999149

<400> 6801
tgtcttggtta caggacaggc acttgtagcg gcggcccag cgctttagc c 51

<210> 6802
<211> 51
<212> DNA
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<221> misc_feature
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<400> 6802
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<210> 6803
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<400> 6803
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<210> 6804
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<210> 6805
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<221> misc_feature
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<223> Accession number cg43273813

<400> 6805
tataaaaaat tagccaggca tggtagcata tgctgtagt cccagctact c 51

<210> 6806
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<221> misc_feature
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<210> 6807
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<223> 1 of 2 allelic variants (6808 is other entry)

<221> misc_feature
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<223> Accession number cg42487874

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gtcaggagtt ttagaccatc ctggccagca cggtagacc cctctctac t 51

<210> 6808
<211> 51

<212> DNA
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<221> misc_feature
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<400> 6808
gtcaggagtt tgagaccatc ctggctagca cggatgaagcc ccgtctctac t

51

<210> 6809
<211> 14
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<213> Homo sapiens

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<222> (7)...(0)
<223> cSNP translation

<221> misc_feature
<222> (0)...(0)
<223> Peptide 1 of 2 allelic variants (6810 is other peptide)

<400> 6809
Ile Ala Leu Gln Glu Ala Arg Asp Ile Cys Glu Gly Gln Val
1 5 10

<210> 6810
<211> 14
<212> PRT
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<222> (7)...(0)
<223> cSNP translation

<221> misc_feature
<222> (0)...(0)
<223> Peptide 2 of 2 allelic variants (6809 is other peptide)
<400> 6810
Ile Ala Leu Gln Glu Ala Lys Asp Ile Cys Glu Gly Gln Val
1 5 10

<210> 6811
<211> 14
<212> PRT
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<222> (7)...(0)
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<221> misc_feature
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<223> Peptide 1 of 2 allelic variants (6812 is other peptide)

<400> 6811
Leu Ala Ala Arg Val Ala Val Leu Arg Asp Gln Gly Val Val
1 5 10

<210> 6812
<211> 14
<212> PRT
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<221> misc_feature
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<223> Peptide 2 of 2 allelic variants (6811 is other peptide)

<400> 6812
Leu Ala Ala Arg Val Ala Ala Leu Arg Asp Gln Gly Val Val
1 5 10

<210> 6813
<211> 14
<212> PRT
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<222> (7)...(0)
<223> cSNP translation

<221> misc_feature
<222> (0)...(0)
<223> Peptide 1 of 2 allelic variants (6814 is other peptide)

<400> 6813
Ala Thr Ser Asp Pro Glu Glu Phe Thr Thr Gly Arg Trp Arg
1 5 10

<210> 6814
<211> 14
<212> PRT
<213> Homo sapiens

<220>
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<222> (7)...(0)
<223> cSNP translation

<221> misc_feature

<222> (0)...(0)
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<400> 6814
Ala Thr Ser Asp Pro Glu Asp Phe Thr Thr Gly Arg Trp Arg
1 5 10

<210> 6815
<211> 14
<212> PRT
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<222> (7)...(0)
<223> cSNP translation

<221> misc_feature
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<223> Peptide 1 of 2 allelic variants (6816 is other peptide)

<400> 6815
Arg Thr Asp Phe Ala Ile Glu Val Cys His Ser Val Met Asp
1 5 10

<210> 6816
<211> 14
<212> PRT
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<222> (7)...(0)
<223> cSNP translation

<221> misc_feature
<222> (0)...(0)
<223> Peptide 2 of 2 allelic variants (6815 is other peptide)
<400> 6816
Arg Thr Asp Phe Ala Ile Asp Val Cys His Ser Val Met Asp
1 5 10

<210> 6817
<211> 14
<212> PRT
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<220>
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<222> (7)...(0)
<223> cSNP translation

<221> misc_feature
<222> (0)...(0)
<223> Peptide 1 of 2 allelic variants (6818 is other peptide)

<400> 6817
Leu Glu Glu Val Asp Ser Val Val Ala Ala Glu Tyr Glu Leu

1 5 10

<210> 6818
<211> 14
<212> PRT
<213> Homo sapiens

<220>
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<223> cSNP translation

<221> misc_feature
<222> (0)...(0)
<223> Peptide 2 of 2 allelic variants (6817 is other peptide)
<400> 6818
Leu Glu Glu Val Asp Ser Ile Val Ala Ala Glu Tyr Glu Leu
1 5 10

<210> 6819
<211> 14
<212> PRT
<213> Homo sapiens

<220>
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<223> cSNP translation

<221> misc_feature
<222> (0)...(0)
<223> Peptide 1 of 2 allelic variants (6820 is other peptide)
<400> 6819
Lys Gly Leu Glu Val Gly Asp Ser Leu Leu Asp Glu Asp Ser
1 5 10

<210> 6820
<211> 14
<212> PRT
<213> Homo sapiens

<220>
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<221> misc_feature
<222> (0)...(0)
<223> Peptide 2 of 2 allelic variants (6819 is other peptide)
<400> 6820
Lys Gly Leu Glu Val Gly Glu Ser Leu Leu Asp Glu Asp Ser
1 5 10

<210> 6821
<211> 14
<212> PRT

<213> Homo sapiens

<220>

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<222> (7)...(0)

<223> cSNP translation

<221> misc_feature

<222> (0)...(0)

<223> Peptide 1 of 2 allelic variants (6822 is other peptide)

<400> 6821

Leu Arg Leu Leu Ala Gly Arg Asp Pro His His Ile Cys Glu
1 5 10

<210> 6822

<211> 14

<212> PRT

<213> Homo sapiens

<220>

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<222> (7)...(0)

<223> cSNP translation

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<222> (0)...(0)

<223> Peptide 2 of 2 allelic variants (6821 is other peptide)

<400> 6822

Leu Arg Leu Leu Ala Gly His Asp Pro His His Ile Cys Glu
1 5 10

<210> 6823

<211> 14

<212> PRT

<213> Homo sapiens

<220>

<221> VARIANT

<222> (7)...(0)

<223> cSNP translation

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<222> (0)...(0)

<223> Peptide 1 of 2 allelic variants (6824 is other peptide)

<400> 6823

Glu Leu Arg His Val Leu Ala Thr Leu Gly Glu Arg Leu Thr
1 5 10

<210> 6824

<211> 14

<212> PRT

<213> Homo sapiens

<220>

<221> VARIANT

<222> (7)...(0)
<223> cSNP translation

<221> misc_feature
<222> (0)...(0)
<223> Peptide 2 of 2 allelic variants (6823 is other peptide)
<400> 6824
Glu Leu Arg His Val Leu Val Thr Leu Gly Glu Arg Leu Thr
1 5 10

<210> 6825
<211> 14
<212> PRT
<213> Homo sapiens

<220>
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<222> (7)...(0)
<223> cSNP translation

<221> misc_feature
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<223> Peptide 1 of 2 allelic variants (6826 is other peptide)

<400> 6825
Glu Asp Arg Met Asp Thr Val Glu Val Leu Lys Arg Asn Gly
1 5 10

<210> 6826
<211> 14
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<223> Peptide 2 of 2 allelic variants (6825 is other peptide)
<400> 6826
Glu Asp Arg Met Asp Thr Ile Glu Val Leu Lys Arg Asn Gly
1 5 10

<210> 6827
<211> 14
<212> PRT
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<223> Peptide 1 of 2 allelic variants (6828 is other peptide)

<400> 6827

Met His Leu Leu Thr His Asp Leu Arg His Lys Cys Gly Val
1 5 10

<210> 6828

<211> 14

<212> PRT

<213> Homo sapiens

<220>

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<223> cSNP translation

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<223> Peptide 2 of 2 allelic variants (6827 is other peptide)

<400> 6828

Met His Leu Leu Thr His Asn Leu Arg His Lys Cys Gly Val
1 5 10

<210> 6829

<211> 14

<212> PRT

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<223> Peptide 1 of 2 allelic variants (6830 is other peptide)

<400> 6829

Ser Ser His Glu Gly Lys Val Ala Ser Tyr Lys His Phe Thr
1 5 10

<210> 6830

<211> 14

<212> PRT

<213> Homo sapiens

<220>

<221> VARIANT

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<223> cSNP translation

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<222> (0)...(0)

<223> Peptide 2 of 2 allelic variants (6829 is other peptide)

<400> 6830

Ser Ser His Glu Gly Lys Ile Ala Ser Tyr Lys His Phe Thr
1 5 10

<210> 6831
<211> 5
<212> PRT
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<221> misc_feature
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<223> Peptide 1 of 2 allelic variants (6832 is other peptide)

<400> 6831
Trp Pro Gly Trp Ser
1 5

<210> 6832
<211> 5
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<221> misc_feature
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<223> Peptide 2 of 2 allelic variants (6831 is other peptide)

<400> 6832
Trp Pro Gly Trp Ser
1 5

<210> 6833
<211> 14
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<220>
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<223> Peptide 1 of 2 allelic variants (6834 is other peptide)

<400> 6833
Leu Thr Ala Ala Ile Ser Lys Val Leu His Asp Lys Tyr Pro
1 5 10

<210> 6834
<211> 14
<212> PRT
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<220>
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<223> Peptide 2 of 2 allelic variants (6833 is other peptide)

<400> 6834

Leu Thr Ala Ala Ile Ser Arg Val Leu His Asp Lys Tyr Pro
1 5 10

<210> 6835

<211> 14

<212> PRT

<213> Homo sapiens

<220>

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<222> (7)...(0)

<223> cSNP translation

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<223> Peptide 1 of 2 allelic variants (6836 is other peptide)

<400> 6835

Ser Ser Trp Asp Tyr Arg His Ala Pro Pro Arg Pro Ala Asn
1 5 10

<210> 6836

<211> 14

<212> PRT

<213> Homo sapiens

<220>

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<223> cSNP translation

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<223> Peptide 2 of 2 allelic variants (6835 is other peptide)

<400> 6836

Ser Ser Trp Asp Tyr Arg Arg Ala Pro Pro Arg Pro Ala Asn
1 5 10

<210> 6837

<211> 14

<212> PRT

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<221> VARIANT

<222> (7)...(0)

<223> cSNP translation

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<222> (0)...(0)

<223> Peptide 1 of 2 allelic variants (6838 is other peptide)

<400> 6837

Tyr Arg Arg Val Pro Pro Arg Leu Ala Asn Phe Val Phe Leu
1 5 10

<210> 6838
<211> 14
<212> PRT
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<223> Peptide 2 of 2 allelic variants (6837 is other peptide)
<400> 6838
Tyr Arg Arg Val Pro Pro His Leu Ala Asn Phe Val Phe Leu
1 5 10

<210> 6839
<211> 14
<212> PRT
<213> Homo sapiens

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<223> Peptide 1 of 2 allelic variants (6840 is other peptide)

<400> 6839
Gly Phe Gly His Ser Ser Asp Phe Lys Arg His Arg Arg Thr
1 5 10

<210> 6840
<211> 14
<212> PRT
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<223> Peptide 2 of 2 allelic variants (6839 is other peptide)
<400> 6840
Gly Phe Gly His Ser Ser Asn Phe Lys Arg His Arg Arg Thr
1 5 10

<210> 6841
<211> 14
<212> PRT
<213> Homo sapiens

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<223> Peptide 1 of 2 allelic variants (6842 is other peptide)

<400> 6841
Pro Gly Trp Asn Pro Gln Ile Pro Glu Lys Lys Gly Lys Glu
1 5 10

<210> 6842
<211> 14
<212> PRT
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<223> Peptide 2 of 2 allelic variants (6841 is other peptide)
<400> 6842

Pro Gly Trp Asn Pro Gln Val Pro Glu Lys Lys Gly Lys Glu
1 5 10

<210> 6843
<211> 14
<212> PRT
<213> Homo sapiens

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<223> Peptide 1 of 2 allelic variants (6844 is other peptide)

<400> 6843
Arg Ser Arg Cys Pro Glu Val Gly Leu Arg Ser Asn Phe Ile
1 5 10

<210> 6844
<211> 14
<212> PRT
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<221> misc_feature
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<223> Peptide 2 of 2 allelic variants (6843 is other peptide)
<400> 6844
Arg Ser Arg Cys Pro Glu Ala Gly Leu Arg Ser Asn Phe Ile
1 5 10

<210> 6845
<211> 14
<212> PRT
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<222> (0)...(0)
<223> Peptide 1 of 2 allelic variants (6846 is other peptide)
<400> 6845
Thr Pro Asp Leu Lys Ile His Leu Pro Arg Pro Pro Lys Val
1 5 10

<210> 6846
<211> 14
<212> PRT
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<222> (0)...(0)
<223> Peptide 2 of 2 allelic variants (6845 is other peptide)
<400> 6846
Thr Pro Asp Leu Lys Ile Arg Leu Pro Arg Pro Pro Lys Val
1 5 10

<210> 6847
<211> 14
<212> PRT
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<223> Peptide 1 of 2 allelic variants (6848 is other peptide)

<400> 6847

Val Trp Asn Met Arg Leu Val Phe Phe Phe Gly Val Ser Ile
1 5 10

<210> 6848

<211> 14

<212> PRT

<213> Homo sapiens

<220>

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<222> (7)...(0)

<223> cSNP translation

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<222> (0)...(0)

<223> Peptide 2 of 2 allelic variants (6847 is other peptide)

<400> 6848

Val Trp Asn Met Arg Leu Ala Phe Phe Phe Gly Val Ser Ile
1 5 10

<210> 6849

<211> 14

<212> PRT

<213> Homo sapiens

<220>

<221> VARIANT

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<223> cSNP translation

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<222> (0)...(0)

<223> Peptide 1 of 2 allelic variants (6850 is other peptide)

<400> 6849

Gln Leu Ser Val Arg Ile Glu Leu Arg His Phe Gln Leu Cys
1 5 10

<210> 6850

<211> 14

<212> PRT

<213> Homo sapiens

<220>

<221> VARIANT

<222> (7)...(0)

<223> cSNP translation

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<222> (0)...(0)

<223> Peptide 2 of 2 allelic variants (6849 is other peptide)

<400> 6850

Gln Leu Ser Val Arg Ile Gln Leu Arg His Phe Gln Leu Cys
1 5 10

<210> 6851

<211> 14
<212> PRT
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<223> Peptide 1 of 2 allelic variants (6852 is other peptide)

<400> 6851
Asn Tyr Gln Leu Ser Val Arg Ile Glu Leu Arg His Phe Gln
1 5 10

<210> 6852
<211> 14
<212> PRT
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<223> Peptide 2 of 2 allelic variants (6851 is other peptide)
<400> 6852

Asn Tyr Gln Leu Ser Val Lys Ile Glu Leu Arg His Phe Gln
1 5 10

<210> 6853
<211> 9
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<400> 6853
Leu Ala Gly Arg Gly Gly Ala Arg Leu
1 5

<210> 6854
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<223> Peptide 2 of 2 allelic variants (6853 is other peptide)
<400> 6854
Leu Ala Gly Arg Gly Gly Gly Arg Leu
1 5

<210> 6855
<211> 14
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<223> Peptide 1 of 2 allelic variants (6856 is other peptide),

<400> 6855
His Cys Thr Pro Ala Trp Ala Thr Glu Gln Asp Ser Val Ser
1 5 10

<210> 6856
<211> 14
<212> PRT
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<400> 6856
His Cys Thr Pro Ala Trp Val Thr Glu Gln Asp Ser Val Ser
1 5 10

<210> 6857
<211> 14
<212> PRT
<213> Homo sapiens

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<223> Peptide 1 of 2 allelic variants (6858 is other peptide)

<400> 6857
Ala His Leu Gly Leu Pro Lys Cys Trp Asp Tyr Arg His Glu
1 5 10

<210> 6858
<211> 14
<212> PRT
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<223> Peptide 2 of 2 allelic variants (6857 is other peptide)
<400> 6858
Ala His Leu Gly Leu Pro Arg Cys Trp Asp Tyr Arg His Glu
1 5 10

<210> 6859
<211> 14
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<223> Peptide 1 of 2 allelic variants (6860 is other peptide)

<400> 6859
Thr Gly Arg Val Ile Gly Gly Val Asp Glu Ala Ala Ala Asp
1 5 10

<210> 6860
<211> 14
<212> PRT
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<223> Peptide 2 of 2 allelic variants (6859 is other peptide)
<400> 6860

Thr Gly Arg Val Ile Gly Val Val Asp Glu Ala Ala Ala Asp
1 5 10

<210> 6861

<211> 3

<212> PRT

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<221> misc_feature

<222> (0)...(0)

<223> Peptide 1 of 2 allelic variants (6862 is other peptide)

<400> 6861

Asp Leu Lys

1

<210> 6862

<211> 3

<212> PRT

<213> Homo sapiens

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<222> (0)...(0)

<223> Peptide 2 of 2 allelic variants (6861 is other peptide)

<400> 6862

Asp Leu Lys

1

<210> 6863

<211> 14

<212> PRT

<213> Homo sapiens

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<222> (0)...(0)

<223> Peptide 1 of 2 allelic variants (6864 is other peptide)

<400> 6863

Cys Phe Ser Arg Leu Ser Thr Leu Leu Glu His Arg His Thr

1

5

10

<210> 6864

<211> 14

<212> PRT

<213> Homo sapiens

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<223> cSNP translation

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<223> Peptide 2 of 2 allelic variants (6863 is other peptide)
<400> 6864

Cys Phe Ser Arg Leu Ser Ser Leu Leu Glu His Arg His Thr
1 5 10

<210> 6865
<211> 14
<212> PRT
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<223> Peptide 1 of 2 allelic variants (6866 is other peptide)

<400> 6865
Asp Glu Leu Ala Glu Val Phe Ala Pro Tyr Thr Asn Val Thr
1 5 10

<210> 6866
<211> 14
<212> PRT
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<400> 6866
Asp Glu Leu Ala Glu Val Leu Ala Pro Tyr Thr Asn Val Thr
1 5 10

<210> 6867
<211> 14
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<400> 6867
Tyr Glu Tyr Ser Pro Glu Ile Phe Ser Gln Thr Arg Thr Asp

1 5 10

<210> 6868
<211> 14
<212> PRT
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<223> Peptide 2 of 2 allelic variants (6867 is other peptide)
<400> 6868
Tyr Glu Tyr Ser Pro Glu Met Phe Ser Gln Thr Arg Thr Asp
1 5 10

<210> 6869
<211> 14
<212> PRT
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<223> Peptide 1 of 2 allelic variants (6870 is other peptide)
<400> 6869
Pro Ala Thr Val Glu Met Gly Thr Pro Asn Thr Tyr Ala Asp
1 5 10

<210> 6870
<211> 14
<212> PRT
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<223> Peptide 2 of 2 allelic variants (6869 is other peptide)
<400> 6870
Pro Ala Thr Val Glu Met Ser Thr Pro Asn Thr Tyr Ala Asp
1 5 10

<210> 6871
<211> 14
<212> PRT

<213> Homo sapiens

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<221> VARIANT

<222> (7)...(0)

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<222> (0)...(0)

<223> Peptide 1 of 2 allelic variants (6872 is other peptide)

<400> 6871

Thr Gly Leu Trp Glu Ser Gly Pro Glu Asp Gln Leu Thr Thr
1 5 10

<210> 6872

<211> 14

<212> PRT

<213> Homo sapiens

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<221> VARIANT

<222> (7)...(0)

<223> cSNP translation

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<222> (0)...(0)

<223> Peptide 2 of 2 allelic variants (6871 is other peptide)

<400> 6872

Thr Gly Leu Trp Glu Ser Cys Pro Glu Asp Gln Leu Thr Thr
1 5 10

<210> 6873

<211> 14

<212> PRT

<213> Homo sapiens

<220>

<221> VARIANT

<222> (7)...(0)

<223> cSNP translation

<221> misc_feature

<222> (0)...(0)

<223> Peptide 1 of 2 allelic variants (6874 is other peptide)

<400> 6873

Pro Glu Ile Ser Phe Thr Ser Asp Ser Ser Phe Ala Lys Gly
1 5 10

<210> 6874

<211> 14

<212> PRT

<213> Homo sapiens

<220>

<221> VARIANT

<222> (7)...(0)
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<223> Peptide 2 of 2 allelic variants (6873 is other peptide)
<400> 6874
Pro Glu Ile Ser Phe Thr Ala Asp Ser Ser Phe Ala Lys Gly
1 5 10

<210> 6875
<211> 8
<212> PRT
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<223> Peptide 1 of 2 allelic variants (6876 is other peptide)

<400> 6875
Gly Leu Asp Ser Tyr Gln Arg Asp
1 5

<210> 6876
<211> 8
<212> PRT
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<223> Peptide 2 of 2 allelic variants (6875 is other peptide)
<400> 6876
Gly Leu Asp Ser Tyr Gln Gly Asp
1 5

<210> 6877
<211> 14
<212> PRT
<213> Homo sapiens

<220>
<221> VARIANT
<222> (7)...(0)
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<221> misc_feature
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<223> Peptide 1 of 2 allelic variants (6878 is other peptide)

<400> 6877

Thr Pro Arg Arg Ser Arg Gly Gln Glu Ile Ile Lys Leu Arg
1 5 10

<210> 6878

<211> 14

<212> PRT

<213> Homo sapiens

<220>

<221> VARIANT

<222> (7)...(0)

<223> cSNP translation

<221> misc_feature

<222> (0)...(0)

<223> Peptide 2 of 2 allelic variants (6877 is other peptide)

<400> 6878

Thr Pro Arg Arg Ser Arg Arg Gln Glu Ile Ile Lys Leu Arg
1 5 10

<210> 6879

<211> 14

<212> PRT

<213> Homo sapiens

<220>

<221> VARIANT

<222> (7)...(0)

<223> cSNP translation

<221> misc_feature

<222> (0)...(0)

<223> Peptide 1 of 2 allelic variants (6880 is other peptide)

<400> 6879

Val Glu Glu Gln Thr Glu Glu Thr Gln Val Thr Glu Glu Val
1 5 10

<210> 6880

<211> 14

<212> PRT

<213> Homo sapiens

<220>

<221> VARIANT

<222> (7)...(0)

<223> cSNP translation

<221> misc_feature

<222> (0)...(0)

<223> Peptide 2 of 2 allelic variants (6879 is other peptide)

<400> 6880

Val Glu Glu Gln Thr Glu Lys Thr Gln Val Thr Glu Glu Val
1 5 10

<210> 6881

<211> 14

<212> PRT

<213> Homo sapiens

<220>

<221> VARIANT

<222> (7)...(0)

<223> cSNP translation

<221> misc_feature

<222> (0)...(0)

<223> Peptide 1 of 2 allelic variants (6882 is other peptide)

<400> 6881

Cys Lys Asp Lys Gln Leu Gln Ile Phe Asp Pro Arg Thr Lys
1 5 10

<210> 6882

<211> 14

<212> PRT

<213> Homo sapiens

<220>

<221> VARIANT

<222> (7)...(0)

<223> cSNP translation

<221> misc_feature

<222> (0)...(0)

<223> Peptide 2 of 2 allelic variants (6881 is other peptide)

<400> 6882

Cys Lys Asp Lys Gln Leu Arg Ile Phe Asp Pro Arg Thr Lys
1 5 10

<210> 6883

<211> 14

<212> PRT

<213> Homo sapiens

<220>

<221> VARIANT

<222> (7)...(0)

<223> cSNP translation

<221> misc_feature

<222> (0)...(0)

<223> Peptide 1 of 2 allelic variants (6884 is other peptide)

<400> 6883

Cys Leu Gln Asp Leu Arg Ser Leu Glu Gly Cys Tyr Pro Lys
1 5 10

<210> 6884

<211> 14

<212> PRT

<213> Homo sapiens

<220>

<221> VARIANT

<222> (7)...(0)

<223> cSNP translation

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<222> (0)...(0)

<223> Peptide 2 of 2 allelic variants (6883 is other peptide)

<400> 6884

Cys Leu Gln Asp Leu Arg Gly Leu Glu Gly Cys Tyr Pro Lys
1 5 10

<210> 6885

<211> 14

<212> PRT

<213> Homo sapiens

<220>

<221> VARIANT

<222> (7)...(0)

<223> cSNP translation

<221> misc_feature

<222> (0)...(0)

<223> Peptide 1 of 2 allelic variants (6886 is other peptide)

<400> 6885

Leu Gly Phe Val Asp Asp Lys Lys Ala Ala Ser Ile Lys Glu
1 5 10

<210> 6886

<211> 14

<212> PRT

<213> Homo sapiens

<220>

<221> VARIANT

<222> (7)...(0)

<223> cSNP translation

<221> misc_feature

<222> (0)...(0)

<223> Peptide 2 of 2 allelic variants (6885 is other peptide)

<400> 6886

Leu Gly Phe Val Asp Asp Glu Lys Ala Ala Ser Ile Lys Glu
1 5 10

<210> 6887

<211> 14

<212> PRT

<213> Homo sapiens

<220>

<221> VARIANT

<222> (7)...(0)

<223> cSNP translation

<221> misc_feature

<222> (0)...(0)

<223> Peptide 1 of 2 allelic variants (6888 is other peptide)

<400> 6887

Ala Arg Ser His Tyr Gly Glu Ile Cys Ser Thr His Ser Tyr
1 5 10

<210> 6888

<211> 14

<212> PRT

<213> Homo sapiens

<220>

<221> VARIANT

<222> (7)...(0)

<223> cSNP translation

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<223> Peptide 2 of 2 allelic variants (6887 is other peptide)

<400> 6888

Ala Arg Ser His Tyr Gly Lys Ile Cys Ser Thr His Ser Tyr
1 5 10

<210> 6889

<211> 14

<212> PRT

<213> Homo sapiens

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<223> Peptide 1 of 2 allelic variants (6890 is other peptide)

<400> 6889

Gly Thr Pro Glu Glu Leu Gln Glu Leu Val Asp Thr Ala His
1 5 10

<210> 6890

<211> 14

<212> PRT

<213> Homo sapiens

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<223> Peptide 2 of 2 allelic variants (6889 is other peptide)

<400> 6890

Gly Thr Pro Glu Glu Leu Lys Glu Leu Val Asp Thr Ala His
1 5 10

<210> 6891

<211> 14

<212> PRT

<213> Homo sapiens

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<223> Peptide 1 of 2 allelic variants (6892 is other peptide)

<400> 6891

Ser Tyr Lys His Phe Thr Cys Asn Val Leu Pro Arg Ile Lys
1 5 10

<210> 6892

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<223> Peptide 2 of 2 allelic variants (6891 is other peptide)

<400> 6892

Ser Tyr Lys His Phe Thr Ser Asn Val Leu Pro Arg Ile Lys
1 5 10

<210> 6893

<211> 14

<212> PRT

<213> Homo sapiens

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<223> Peptide 1 of 2 allelic variants (6894 is other peptide)

<400> 6893

Tyr Glu Phe Lys His Ser Arg Pro Lys Lys Pro Arg Ser Leu
1 5 10

<210> 6894
<211> 14
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<400> 6894
Tyr Glu Phe Lys His Ser Gly Pro Lys Lys Pro Arg Ser Leu
1 5 10

<210> 6895
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<400> 6895
Pro Pro Ile Thr Gln Leu Gly Asn Tyr Ala Gly Arg Tyr Ile
1 5 10

<210> 6896
<211> 14
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<223> cSNP translation

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<223> Peptide 2 of 2 allelic variants (6895 is other peptide)
<400> 6896
Pro Pro Ile Thr Gln Leu Asp Asn Tyr Ala Gly Arg Tyr Ile
1 5 10

<210> 6897
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<400> 6897
Arg Asp Pro Asp Met Glu Met Ile Ala Arg Ala Arg Ile Leu
1 5 10

<210> 6898
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<400> 6898
Arg Asp Pro Asp Met Glu Thr Ile Ala Arg Ala Arg Ile Leu
1 5 10

<210> 6899
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<400> 6899
Arg Val Leu Ala Gly Glu Thr Leu Pro Ala Ala Gly Ser Val
1 5 10

<210> 6900
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<212> PRT
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<223> Peptide 2 of 2 allelic variants (6899 is other peptide)

<400> 6900

Arg Val Leu Ala Gly Glu Ala Leu Pro Ala Ala Gly Ser Val
1 5 10

<210> 6901

<211> 14

<212> PRT

<213> Homo sapiens

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<223> Peptide 1 of 2 allelic variants (6902 is other peptide)

<400> 6901

Ser Thr Thr Asp Leu Arg Val Asp Tyr Lys Tyr Asn Pro Glu
1 5 10

<210> 6902

<211> 14

<212> PRT

<213> Homo sapiens

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<223> Peptide 2 of 2 allelic variants (6901 is other peptide)

<400> 6902

Ser Thr Thr Asp Leu Arg Met Asp Tyr Lys Tyr Asn Pro Glu
1 5 10

<210> 6903

<211> 14

<212> PRT

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<223> cSNP translation

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<222> (0)...(0)

<223> Peptide 1 of 2 allelic variants (6904 is other peptide)

<400> 6903
Ala Arg Ser Val Leu Asp Gly Asp Ser Val Arg Ala Arg Trp
1 5 10

<210> 6904
<211> 14
<212> PRT
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<223> Peptide 2 of 2 allelic variants (6903 is other peptide)

<400> 6904
Ala Arg Ser Val Leu Asp Trp Asp Ser Val Arg Ala Arg Trp
1 5 10

<210> 6905
<211> 14
<212> PRT
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<223> Peptide 1 of 2 allelic variants (6906 is other peptide)

<400> 6905
Asn Lys Glu His Ser Phe Glu Val Ser Leu Phe Ala Glu Leu
1 5 10

<210> 6906
<211> 6
<212> PRT
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<221> misc_feature
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<223> Peptide 2 of 2 allelic variants (6905 is other peptide)

<400> 6906
Asn Lys Glu His Ser Phe
1 5

<210> 6907
<211> 4
<212> PRT
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<221> misc_feature
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<223> Peptide 1 of 2 allelic variants (6908 is other peptide)

<400> 6907
Pro Asp Leu Lys
1

<210> 6908
<211> 14
<212> PRT
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<221> misc_feature
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<223> Peptide 2 of 2 allelic variants (6907 is other peptide)
<400> 6908
Pro Asp Leu Lys
1

<210> 6909
<211> 14
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<223> Peptide 1 of 2 allelic variants (6910 is other peptide)

<400> 6909
Glu Asp Gly Phe Leu Gln Gly Phe Lys Gly Gln Leu Ile His
1 5 10

<210> 6910
<211> 14
<212> PRT
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<223> Peptide 2 of 2 allelic variants (6909 is other peptide)
<400> 6910
Glu Asp Gly Phe Leu Gln Arg Phe Lys Gly Gln Leu Ile His
1 5 10

<210> 6911
<211> 14

<212> PRT
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<400> 6911
Thr Glu Asp Lys Thr Val Arg Glu Asn Val Glu Glu Ala Val
1 5 10

<210> 6912
<211> 14
<212> PRT
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<223> Peptide 2 of 2 allelic variants (6911 is other peptide)

<400> 6912
Thr Glu Asp Lys Thr Val Cys Glu Asn Val Glu Glu Ala Val
1 5 10

<210> 6913
<211> 1
<212> PRT
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<221> misc_feature
<222> (0)...(0)
<223> Peptide 1 of 2 allelic variants (6914 is other peptide)

<400> 6913
His
1

<210> 6914
<211> 1
<212> PRT
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<221> misc_feature
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<223> Peptide 2 of 2 allelic variants (6913 is other peptide)

<400> 6914
His
1

<210> 6915
<211> 11
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<223> Peptide 1 of 2 allelic variants (6916 is other peptide)

<400> 6915
Pro Arg Ser Cys His Cys Ala Pro Ala Trp Ala
1 5 10

<210> 6916
<211> 11
<212> PRT
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<223> Peptide 2 of 2 allelic variants (6915 is other peptide)

<400> 6916
Pro Arg Ser Cys His Cys Thr Pro Ala Trp Ala
1 5 10

<210> 6917
<211> 14
<212> PRT
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<223> Peptide 1 of 2 allelic variants (6918 is other peptide)

<400> 6917
Pro Gln Asp Phe Ile Gly Cys Leu Asn Val Lys Ala Thr Phe
1 5 10

<210> 6918
<211> 14
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<222> (7)...(0)

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<223> Peptide 2 of 2 allelic variants (6917 is other peptide)

<400> 6918

Pro Gln Asp Phe Ile Gly Ser Leu Asn Val Lys Ala Thr Phe

1 5 10

<210> 6919

<211> 14

<212> PRT

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<223> Peptide 1 of 2 allelic variants (6920 is other peptide)

<400> 6919

Thr Thr Pro Leu Ser Asp Asp His Asp Glu Lys Tyr Gly Val

1 5 10

<210> 6920

<211> 14

<212> PRT

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<223> Peptide 2 of 2 allelic variants (6919 is other peptide)

<400> 6920

Thr Thr Pro Leu Ser Asp Gly His Asp Glu Lys Tyr Gly Val

1 5 10

<210> 6921

<211> 14

<212> PRT

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<222> (0)...(0)

<223> Peptide 1 of 2 allelic variants (6922 is other peptide)

<400> 6921

Gln Ala Gly Val Gln Trp Cys Asn Leu Gly Ser Leu Gln Pro
1 5 10

<210> 6922

<211> 14

<212> PRT

<213> Homo sapiens

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<223> Peptide 2 of 2 allelic variants (6921 is other peptide)

<400> 6922

Gln Ala Gly Val Gln Trp Arg Asn Leu Gly Ser Leu Gln Pro
1 5 10

<210> 6923

<211> 14

<212> PRT

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<223> Peptide 1 of 2 allelic variants (6924 is other peptide)

<400> 6923

Ser Trp Asp Tyr Arg His Ala Pro Pro Arg Pro Ala Asn Phe
1 5 10

<210> 6924

<211> 14

<212> PRT

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<223> Peptide 2 of 2 allelic variants (6923 is other peptide)

<400> 6924

Ser Trp Asp Tyr Arg His Pro Pro Pro Arg Pro Ala Asn Phe
1 5 10

<210> 6925

<211> 13

<212> PRT

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<223> Peptide 1 of 2 allelic variants (6926 is other peptide)

<400> 6925

Tyr Arg His Ala Pro Pro Arg Pro Ala Asn Phe Leu Tyr
1 5 10

<210> 6926

<211> 13

<212> PRT

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<223> Peptide 2 of 2 allelic variants (6925 is other peptide)

<400> 6926

Tyr Arg His Ala Pro Pro Cys Pro Ala Asn Phe Leu Tyr
1 5 10

<210> 6927

<211> 14

<212> PRT

<213> Homo sapiens

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<223> Peptide 1 of 2 allelic variants (6928 is other peptide)

<400> 6927

Phe Cys Ile Phe Ser Arg Asp Gly Val Ser Pro Cys Trp Pro
1 5 10

<210> 6928
<211> 14
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<223> Peptide 2 of 2 allelic variants (6927 is other peptide)
<400> 6928
Phe Cys Ile Phe Ser Arg Gly Gly Val Ser Pro Cys Trp Pro
1 5 10

<210> 6929
<211> 14
<212> PRT
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<223> Peptide 1 of 2 allelic variants (6930 is other peptide)

<400> 6929
Val Ile Gly Trp Lys Lys Ser Glu Gly Ser Pro Pro Pro Glu
1 5 10

<210> 6930
<211> 14
<212> PRT
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<223> Peptide 2 of 2 allelic variants (6929 is other peptide)
<400> 6930
Val Ile Gly Trp Lys Lys Leu Glu Gly Ser Pro Pro Pro Glu
1 5 10

<210> 6931
<211> 13
<212> PRT
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<223> Peptide 1 of 2 allelic variants (6932 is other peptide)

<400> 6931
Tyr Arg His Ala Pro Pro Arg Pro Ala Asn Phe Leu Tyr
1 5 10

<210> 6932
<211> 13
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<223> Peptide 2 of 2 allelic variants (6931 is other peptide)
<400> 6932
Tyr Arg His Ala Pro Pro Cys Pro Ala Asn Phe Leu Tyr
1 5 10

<210> 6933
<211> 14
<212> PRT
<213> Homo sapiens

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<223> Peptide 1 of 2 allelic variants (6934 is other peptide)

<400> 6933
Arg Arg Val Pro Pro Arg Leu Ala Asn Phe Val Phe Leu Val
1 5 10

<210> 6934
<211> 14
<212> PRT
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<223> Peptide 2 of 2 allelic variants (6933 is other peptide)

<400> 6934

Arg Arg Val Pro Pro Arg Pro Ala Asn Phe Val Phe Leu Val
1 5 10

<210> 6935

<211> 9

<212> PRT

<213> Homo sapiens

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<223> Peptide 1 of 2 allelic variants (6936 is other peptide)

<400> 6935

Cys His Pro Gly Trp Ser Thr Val Val
1 5

<210> 6936

<211> 9

<212> PRT

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<222> (7)...(0)

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<223> Peptide 2 of 2 allelic variants (6935 is other peptide)

<400> 6936

Cys His Pro Gly Trp Ser Ala Val Val
1 5

<210> 6937

<211> 14

<212> PRT

<213> Homo sapiens

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<221> VARIANT

<222> (7)...(0)

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<222> (0)...(0)

<223> Peptide 1 of 2 allelic variants (6938 is other peptide)

<400> 6937

Val Thr Ile Ala Cys Gly Ala Leu Ser Gly Phe His Ala Thr
1 5 10

<210> 6938

<211> 14

<212> PRT

<213> Homo sapiens

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<223> cSNP translation

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<222> (0)...(0)

<223> Peptide 2 of 2 allelic variants (6937 is other peptide)

<400> 6938

Val Thr Ile Ala Cys Gly Thr Leu Ser Gly Phe His Ala Thr
1 5 10

<210> 6939

<211> 14

<212> PRT

<213> Homo sapiens

<220>

<221> VARIANT

<222> (7)...(0)

<223> cSNP translation

<221> misc_feature

<222> (0)...(0)

<223> Peptide 1 of 2 allelic variants (6940 is other peptide)

<400> 6939

Leu Leu Gly Arg Leu Arg Gln Glu Asn Cys Leu Asn Pro Gly
1 5 10

<210> 6940

<211> 14

<212> PRT

<213> Homo sapiens

<220>

<221> VARIANT

<222> (7)...(0)

<223> cSNP translation

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<222> (0)...(0)

<223> Peptide 2 of 2 allelic variants (6939 is other peptide)

<400> 6940

Leu Leu Gly Arg Leu Arg Arg Glu Asn Cys Leu Asn Pro Gly
1 5 10

<210> 6941
<211> 14
<212> PRT
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<220>
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<222> (7)...(0)
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<221> misc_feature
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<223> Peptide 1 of 2 allelic variants (6942 is other peptide)

<400> 6941
Gly Asp Ala Glu Ser Phe Leu Asn Ile Ile Asp Ser Ile Arg
1 5 10

<210> 6942
<211> 14
<212> PRT
<213> Homo sapiens

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<222> (7)...(0)
<223> cSNP translation

<221> misc_feature
<222> (0)...(0)
<223> Peptide 2 of 2 allelic variants (6941 is other peptide)

<400> 6942
Gly Asp Ala Glu Ser Phe Pro Asn Ile Ile Asp Ser Ile Arg
1 5 10

<210> 6943
<211> 14
<212> PRT
<213> Homo sapiens

<220>
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<222> (7)...(0)
<223> cSNP translation

<221> misc_feature
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<223> Peptide 1 of 2 allelic variants (6944 is other peptide)

<400> 6943
Tyr Ile Glu Lys Glu Val Lys Tyr Leu Gly Gln Leu Thr Ser
1 5 10

<210> 6944
<211> 14
<212> PRT
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<220>
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<222> (7)...(0)
<223> cSNP translation

<221> misc_feature
<222> (0)...(0)
<223> Peptide 2 of 2 allelic variants (6943 is other peptide)
<400> 6944
Tyr Ile Glu Lys Glu Val Glu Tyr Leu Gly Gln Leu Thr Ser
1 5 10

<210> 6945
<211> 14
<212> PRT
<213> Homo sapiens

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<221> misc_feature
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<223> Peptide 1 of 2 allelic variants (6946 is other peptide)

<400> 6945
Arg Gly Asn Leu Asp Val Ala Lys Leu Asn Gly Asp Trp Phe
1 5 10

<210> 6946
<211> 14
<212> PRT
<213> Homo sapiens

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<221> misc_feature
<222> (0)...(0)
<223> Peptide 2 of 2 allelic variants (6945 is other peptide)
<400> 6946
Arg Gly Asn Leu Asp Val Asp Lys Leu Asn Gly Asp Trp Phe
1 5 10

<210> 6947
<211> 14
<212> PRT
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<223> Peptide 1 of 2 allelic variants (6948 is other peptide)

<400> 6947
Gln Ala Gly Val Gln Trp Arg Asp Leu Ser Ser Leu Gln Thr
1 5 10

<210> 6948
<211> 14
<212> PRT
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<223> Peptide 2 of 2 allelic variants (6947 is other peptide)
<400> 6948
Gln Ala Gly Val Gln Trp Cys Asp Leu Ser Ser Leu Gln Thr
1 5 10

<210> 6949
<211> 13
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<223> Peptide 1 of 2 allelic variants (6950 is other peptide)

<400> 6949
Leu Asn Tyr Pro Ile Leu Glu Gly Trp Phe Val Cys Leu
1 5 10

<210> 6950
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<223> Peptide 2 of 2 allelic variants (6949 is other peptide)

<400> 6950

Leu Asn Tyr Pro Ile Leu Lys Gly Trp Phe Val Cys Leu
1 5 10

<210> 6951

<211> 14

<212> PRT

<213> Homo sapiens

<220>

<221> VARIANT

<222> (7)...(0)

<223> cSNP translation

<221> misc_feature

<222> (0)...(0)

<223> Peptide 1 of 2 allelic variants (6952 is other peptide)

<400> 6951

His Phe Leu Leu Glu Pro Glu Asp Ala Tyr Ile Val Lys Asn
1 5 10

<210> 6952

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<223> Peptide 2 of 2 allelic variants (6951 is other peptide)

<400> 6952

His Phe Leu Leu Glu Pro Gly Asp Ala Tyr Ile Val Lys Asn
1 5 10

<210> 6953

<211> 14

<212> PRT

<213> Homo sapiens

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<223> cSNP translation

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<223> Peptide 1 of 2 allelic variants (6954 is other peptide)

<400> 6953

Leu Ser Ala Arg Val Asp Ala Val Lys Glu Glu Asn Leu Lys
1 5 10

<210> 6954

<211> 14

<212> PRT

<213> Homo sapiens

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<223> cSNP translation

<221> misc_feature

<222> (0)...(0)

<223> Peptide 2 of 2 allelic variants (6953 is other peptide)

<400> 6954

Leu Ser Ala Arg Val Asp Glu Val Lys Glu Glu Asn Leu Lys
1 5 10

<210> 6955

<211> 9

<212> PRT

<213> Homo sapiens

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<223> cSNP translation

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<223> Peptide 1 of 2 allelic variants (6956 is other peptide)

<400> 6955

His Phe Leu Thr Pro Tyr Tyr Lys Asn
1 5

<210> 6956

<211> 6

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<213> Homo sapiens

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<222> (0)...(0)

<223> Peptide 2 of 2 allelic variants (6955 is other peptide)

<400> 6956

His Phe Leu Thr Pro Tyr
1 5

<210> 6957

<211> 6

<212> PRT

<213> Homo sapiens

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<223> Peptide 1 of 2 allelic variants (6958 is other peptide)

<400> 6957
Thr Pro Tyr Tyr Lys Asn
1 5

<210> 6958
<211> 11
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<223> cSNP translation

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<223> Peptide 2 of 2 allelic variants (6957 is other peptide)
<400> 6958
Thr Pro Tyr Tyr Lys Asn Gln Leu Lys Met Asp
1 5 10

<210> 6959
<211> 14
<212> PRT
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<221> misc_feature
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<400> 6959
Arg Phe His Asp Lys Asn Thr Lys Ser Asn Cys Asn Lys Ser
1 5 10

<210> 6960
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<223> Peptide 2 of 2 allelic variants (6959 is other peptide)
<400> 6960
Arg Phe His Asp Lys Asn Ile Lys Ser Asn Cys Asn Lys Ser
1 5 10

<210> 6961
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<221> misc_feature
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<400> 6961
Val Thr Pro Trp Arg His Arg Ile Asn Asn His Ile Gln Thr
1 5 10

<210> 6962
<211> 6
<212> PRT
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<223> Peptide 2 of 2 allelic variants (6961 is other peptide)
<400> 6962
Val Thr Pro Trp Arg His
1 5

<210> 6963
<211> 14
<212> PRT
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<222> (7)...(0)
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<221> misc_feature
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<223> Peptide 1 of 2 allelic variants (6964 is other peptide)

<400> 6963

His Phe Gly Arg Pro Arg Arg Ala Asp His Leu Arg Ser Gly
1 5 10

<210> 6964

<211> 14

<212> PRT

<213> Homo sapiens

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<223> cSNP translation

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<222> (0)...(0)

<223> Peptide 2 of 2 allelic variants (6963 is other peptide)

<400> 6964

His Phe Gly Arg Pro Arg Trp Ala Asp His Leu Arg Ser Gly
1 5 10

<210> 6965

<211> 14

<212> PRT

<213> Homo sapiens

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<222> (7)...(0)

<223> cSNP translation

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<222> (0)...(0)

<223> Peptide 1 of 2 allelic variants (6966 is other peptide)

<400> 6965

Gly Val Asn Pro Gly Gly Gly Ala Cys Ser Glu Pro Ile Ser
1 5 10

<210> 6966

<211> 14

<212> PRT

<213> Homo sapiens

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<223> cSNP translation

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<223> Peptide 2 of 2 allelic variants (6965 is other peptide)

<400> 6966

Gly Val Asn Pro Gly Gly Arg Ala Cys Ser Glu Pro Ile Ser
1 5 10

<210> 6967

<211> 14
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<400> 6967
Phe Val Val Glu Met Gly Phe Arg Tyr Ile Gly Gln Ala Gly
1 5 10

<210> 6968
<211> 14
<212> PRT
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<223> Peptide 2 of 2 allelic variants (6967 is other peptide)

<400> 6968
Phe Val Val Glu Met Gly Leu Arg Tyr Ile Gly Gln Ala Gly
1 5 10

<210> 6969
<211> 14
<212> PRT
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<400> 6969
Glu Ala Glu Ala Gly Gly Ser Leu Lys Val Arg Ser Ser Arg
1 5 10

<210> 6970
<211> 14
<212> PRT
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<223> Peptide 2 of 2 allelic variants (6969 is other peptide)
<400> 6970
Glu Ala Glu Ala Gly Gly Leu Leu Lys Val Arg Ser Ser Arg
1 5 10

<210> 6971
<211> 14
<212> PRT
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<223> Peptide 1 of 2 allelic variants (6972 is other peptide)

<400> 6971
Gly Ser Gly Gly Cys Ser Glu Pro Gly Ser Cys His Cys Thr
1 5 10

<210> 6972
<211> 14
<212> PRT
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<223> Peptide 2 of 2 allelic variants (6971 is other peptide)
<400> 6972
Gly Ser Gly Gly Cys Ser Gly Pro Gly Ser Cys His Cys Thr
1 5 10

<210> 6973
<211> 14
<212> PRT
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<223> Peptide 1 of 2 allelic variants (6974 is other peptide)

<400> 6973
Ala Asn Phe Cys Ile Phe Asn Arg Asp Gly Val Ser Pro Cys
1 5 10

<210> 6974
<211> 14
<212> PRT
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<223> cSNP translation

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<223> Peptide 2 of 2 allelic variants (6973 is other peptide)
<400> 6974
Ala Asn Phe Cys Ile Phe Ser Arg Asp Gly Val Ser Pro Cys
1 5 10

<210> 6975
<211> 14
<212> PRT
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<223> Peptide 1 of 2 allelic variants (6976 is other peptide)

<400> 6975
Lys Arg Phe Ser Cys Leu Ser Leu Leu Ser Ser Trp Asp Tyr
1 5 10

<210> 6976
<211> 14
<212> PRT
<213> Homo sapiens

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<223> cSNP translation

<221> misc_feature
<222> (0)...(0)
<223> Peptide 2 of 2 allelic variants (6975 is other peptide)
<400> 6976

Lys Arg Phe Ser Cys Leu Gly Leu Leu Ser Ser Trp Asp Tyr
1 5 10

<210> 6977

<211> 14

<212> PRT

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<221> VARIANT

<222> (7)...(0)

<223> cSNP translation

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<222> (0)...(0)

<223> Peptide 1 of 2 allelic variants (6978 is other peptide)

<400> 6977

Ala Gly Arg Phe Leu Gly Ser Arg Val Gly Leu Thr Met Asp
1 5 10

<210> 6978

<211> 14

<212> PRT

<213> Homo sapiens

<220>

<221> VARIANT

<222> (7)...(0)

<223> cSNP translation

<221> misc_feature

<222> (0)...(0)

<223> Peptide 2 of 2 allelic variants (6977 is other peptide)

<400> 6978

Ala Gly Arg Phe Leu Gly Gly Arg Val Gly Leu Thr Met Asp
1 5 10

<210> 6979

<211> 14

<212> PRT

<213> Homo sapiens

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<222> (7)...(0)

<223> cSNP translation

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<222> (0)...(0)

<223> Peptide 1 of 2 allelic variants (6980 is other peptide)

<400> 6979

Arg Cys Pro Gln Pro Arg Leu Ala Asn Phe Cys Ile Phe Ser
1 5 10

<210> 6980

<211> 14
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<223> Peptide 2 of 2 allelic variants (6979 is other peptide)
<400> 6980
Arg Cys Pro Gln Pro Arg Pro Ala Asn Phe Cys Ile Phe Ser
1 5 10

<210> 6981
<211> 14
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<223> Peptide 1 of 2 allelic variants (6982 is other peptide)

<400> 6981
Gly Leu Ala His Ser Asp Leu Met Cys Thr Gly Asp Lys Asp
1 5 10

<210> 6982
<211> 14
<212> PRT
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<223> Peptide 2 of 2 allelic variants (6981 is other peptide)
<400> 6982
Gly Leu Ala His Ser Asp Ser Met Cys Thr Gly Asp Lys Asp
1 5 10

<210> 6983
<211> 14
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<223> Peptide 1 of 2 allelic variants (6984 is other peptide)

<400> 6983
Asp Gln Pro Gly Gln His Asp Glu Thr Leu Ser Leu Leu Lys
1 5 10

<210> 6984
<211> 14
<212> PRT
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<221> misc_feature
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<223> Peptide 2 of 2 allelic variants (6983 is other peptide)

<400> 6984
Asp Gln Pro Gly Gln His Gly Glu Thr Leu Ser Leu Leu Lys
1 5 10

<210> 6985
<211> 1
<212> PRT
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<221> misc_feature
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<223> Peptide 1 of 2 allelic variants (6986 is other peptide)

<400> 6985
Leu
1

<210> 6986
<211> 1
<212> PRT
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<221> misc_feature
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<223> Peptide 2 of 2 allelic variants (6985 is other peptide)

<400> 6986
Leu
1

<210> 6987
<211> 14
<212> PRT

<213> Homo sapiens

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<223> cSNP translation

<221> misc_feature

<222> (0)...(0)

<223> Peptide 1 of 2 allelic variants (6988 is other peptide)

<400> 6987

Pro Ala Pro Val Thr Ile Tyr Glu Asn Lys Arg Gln Gly Met
1 5 10

<210> 6988

<211> 14

<212> PRT

<213> Homo sapiens

<220>

<221> VARIANT

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<223> cSNP translation

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<223> Peptide 2 of 2 allelic variants (6987 is other peptide)

<400> 6988

Pro Ala Pro Val Thr Ile His Glu Asn Lys Arg Gln Gly Met
1 5 10

<210> 6989

<211> 14

<212> PRT

<213> Homo sapiens

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<221> VARIANT

<222> (7)...(0)

<223> cSNP translation

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<223> Peptide 1 of 2 allelic variants (6990 is other peptide)

<400> 6989

Asp Tyr Arg Arg Glu Ser Leu Arg Pro Ala Pro Val Thr Ile
1 5 10

<210> 6990

<211> 14

<212> PRT

<213> Homo sapiens

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<223> Peptide 2 of 2 allelic variants (6989 is other peptide)
<400> 6990
Asp Tyr Arg Arg Glu Ser Pro Arg Pro Ala Pro Val Thr Ile
1 5 10

<210> 6991
<211> 14
<212> PRT
<213> Homo sapiens

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<400> 6991
Ser Pro Ala Ser Ala Ser Gln Val Ala Gly Ala Thr Gly Ala
1 5 10

<210> 6992
<211> 14
<212> PRT
<213> Homo sapiens

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<223> Peptide 2 of 2 allelic variants (6991 is other peptide)
<400> 6992
Ser Pro Ala Ser Ala Ser Arg Val Ala Gly Ala Thr Gly Ala
1 5 10

<210> 6993
<211> 14
<212> PRT
<213> Homo sapiens

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<221> misc_feature
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<223> Peptide 1 of 2 allelic variants (6994 is other peptide)

<400> 6993

Arg His Cys His Ser Pro Ala Ser Ala Ser Gln Val Ala Gly
1 5 10

<210> 6994

<211> 14

<212> PRT

<213> Homo sapiens

<220>

<221> VARIANT

<222> (7)...(0)

<223> cSNP translation

<221> misc_feature

<222> (0)...(0)

<223> Peptide 2 of 2 allelic variants (6993 is other peptide)

<400> 6994

Arg His Cys His Ser Pro Thr Ser Ala Ser Gln Val Ala Gly
1 5 10

<210> 6995

<211> 14

<212> PRT

<213> Homo sapiens

<220>

<221> VARIANT

<222> (7)...(0)

<223> cSNP translation

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<222> (0)...(0)

<223> Peptide 1 of 2 allelic variants (6996 is other peptide)

<400> 6995

Ala His Leu Gly Leu Pro Lys Cys Trp Asp Tyr Arg His Glu
1 5 10

<210> 6996

<211> 14

<212> PRT

<213> Homo sapiens

<220>

<221> VARIANT

<222> (7)...(0)

<223> cSNP translation

<221> misc_feature

<222> (0)...(0)

<223> Peptide 2 of 2 allelic variants (6995 is other peptide)

<400> 6996

Ala His Leu Gly Leu Pro Glu Cys Trp Asp Tyr Arg His Glu
1 5 10

<210> 6997
<211> 2
<212> PRT
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<223> Peptide 1 of 2 allelic variants (6998 is other peptide)

<400> 6997
Leu Tyr
1

<210> 6998
<211> 2
<212> PRT
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<223> Peptide 2 of 2 allelic variants (6997 is other peptide)
<400> 6998
Leu Tyr
1

<210> 6999
<211> 14
<212> PRT
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<221> misc_feature
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<223> Peptide 1 of 2 allelic variants (7000 is other peptide)

<400> 6999
Ser Leu Glu Val Arg Ser Ser Arg Pro Val Trp Pro Thr Trp
1 5 10

<210> 7000
<211> 14
<212> PRT

<213> Homo sapiens

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<223> Peptide 2 of 2 allelic variants (6999 is other peptide)

<400> 7000

Ser Leu Glu Val Arg Ser Leu Arg Pro Val Trp Pro Thr Trp

1

5

10

<210> 7001

<211> 14

<212> PRT

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<223> cSNP translation

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<223> Peptide 1 of 2 allelic variants (7002 is other peptide)

<400> 7001

Phe Asp Leu Ala Ile Ala Ala Ala Thr Leu Ala Ala Asp Gly

1

5

10

<210> 7002

<211> 14

<212> PRT

<213> Homo sapiens

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<222> (7)...(0)

<223> cSNP translation

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<223> Peptide 2 of 2 allelic variants (7001 is other peptide)

<400> 7002

Phe Asp Leu Ala Ile Ala Thr Ala Thr Leu Ala Ala Asp Gly

1

5

10

<210> 7003

<211> 14

<212> PRT

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<221> VARIANT

<222> (7)...(0)

<223> cSNP translation

<221> misc_feature

<222> (0)...(0)

<223> Peptide 1 of 2 allelic variants (7004 is other peptide)

<400> 7003

Val Pro Gln Ala Leu Leu Gly Ser Thr Val Leu Ile Gly Glu

1

5

10

<210> 7004

<211> 14

<212> PRT

<213> Homo sapiens

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<222> (7)...(0)

<223> cSNP translation

<221> misc_feature

<222> (0)...(0)

<223> Peptide 2 of 2 allelic variants (7003 is other peptide)

<400> 7004

Val Pro Gln Ala Leu Leu Arg Ser Thr Val Leu Ile Gly Glu

1

5

10

<210> 7005

<211> 7

<212> PRT

<213> Homo sapiens

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<221> VARIANT

<222> (7)...(0)

<223> cSNP translation

<221> misc_feature

<222> (0)...(0)

<223> Peptide 1 of 2 allelic variants (7006 is other peptide)

<400> 7005

Ser Arg Thr Pro Glu Leu Lys

1

5

<210> 7006

<211> 7

<212> PRT

<213> Homo sapiens

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<221> VARIANT

<222> (7)...(0)

<223> cSNP translation

<221> misc_feature

<222> (0)...(0)

<223> Peptide 2 of 2 allelic variants (7005 is other peptide)

<400> 7006

Ser Arg Thr Pro Glu Leu Glu

1

5

<210> 7007

<211> 2

<212> PRT

<213> Homo sapiens

<221> misc_feature

<222> (0)...(0)

<223> Peptide 1 of 2 allelic variants (7008 is other peptide)

<400> 7007

Leu Lys

1

<210> 7008

<211> 2

<212> PRT

<213> Homo sapiens

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<221> VARIANT

<222> (7)...(0)

<223> cSNP translation

<221> misc_feature

<222> (0)...(0)

<223> Peptide 2 of 2 allelic variants (7007 is other peptide)

<400> 7008

Leu Lys

1

<210> 7009

<211> 5

<212> PRT

<213> Homo sapiens

<221> misc_feature

<222> (0)...(0)

<223> Peptide 1 of 2 allelic variants (7010 is other peptide)

<400> 7009

Thr Pro Asp Leu Lys

1

5

<210> 7010

<211> 5

<212> PRT

<213> Homo sapiens

<221> misc_feature

<222> (0)...(0)

<223> Peptide 2 of 2 allelic variants (7009 is other peptide)

<400> 7010

Thr Pro Asp Leu Lys
1 5

<210> 7011

<211> 8

<212> PRT

<213> Homo sapiens

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<221> VARIANT

<222> (7)...(0)

<223> cSNP translation

<221> misc_feature

<222> (0)...(0)

<223> Peptide 1 of 2 allelic variants (7012 is other peptide)

<400> 7011

Ser Pro Phe Trp Leu Gly Trp Ser
1 5

<210> 7012

<211> 6

<212> PRT

<213> Homo sapiens

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<221> VARIANT

<222> (7)...(0)

<223> cSNP translation

<221> misc_feature

<222> (0)...(0)

<223> Peptide 2 of 2 allelic variants (7011 is other peptide)

<400> 7012

Ser Pro Phe Trp Leu Gly
1 5

<210> 7013

<211> 14

<212> PRT

<213> Homo sapiens

<220>

<221> VARIANT

<222> (7)...(0)

<223> cSNP translation

<221> misc_feature

<222> (0)...(0)

<223> Peptide 1 of 2 allelic variants (7014 is other peptide)

<400> 7013

Ser Arg Val Ala Glu Ile Thr Gly Met Arg His His Ala Trp
1 5 10

<210> 7014

<211> 14
<212> PRT
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<223> Peptide 2 of 2 allelic variants (7013 is other peptide)
<400> 7014
Ser Arg Val Ala Glu Ile Ala Gly Met Arg His His Ala Trp
1 5 10

<210> 7015
<211> 14
<212> PRT
<213> Homo sapiens

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<223> Peptide 1 of 2 allelic variants (7016 is other peptide)

<400> 7015
Leu Pro Arg Leu Ser Ala Met Ala Gln Ser Arg Leu His Cys
1 5 10

<210> 7016
<211> 14
<212> PRT
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1 5 10

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Lys Arg Ser Gly Arg Arg Tyr Lys Cys Leu Ser Cys Thr Lys
1 5 10

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Lys Arg Ser Gly Arg Arg Cys Lys Cys Leu Ser Cys Thr Lys
1 5 10

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Phe Trp Trp Pro Ser Leu Leu Met Trp Val Pro Ser Leu Val
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Phe Trp Trp Pro Ser Leu Ser Met Trp Val Pro Ser Leu Val
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Val
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